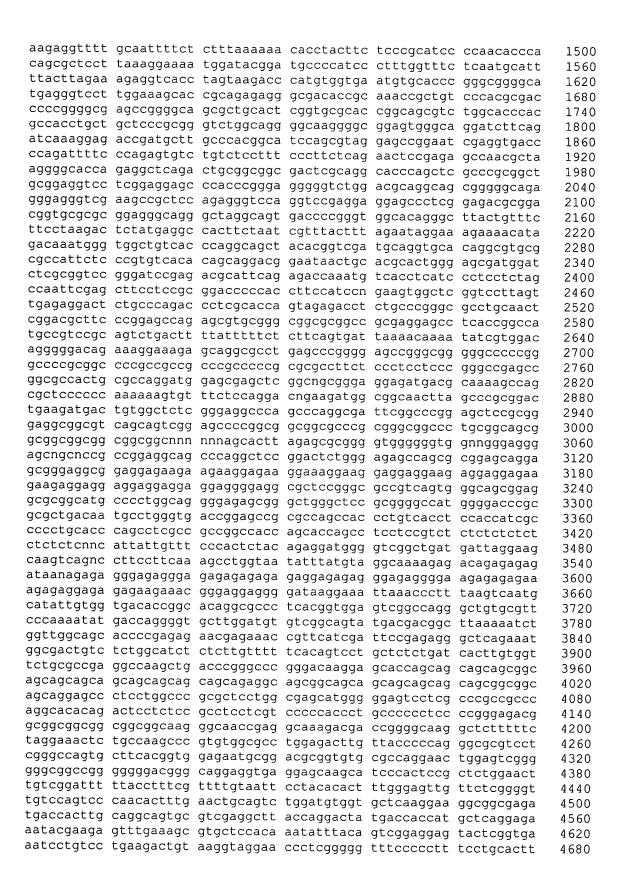


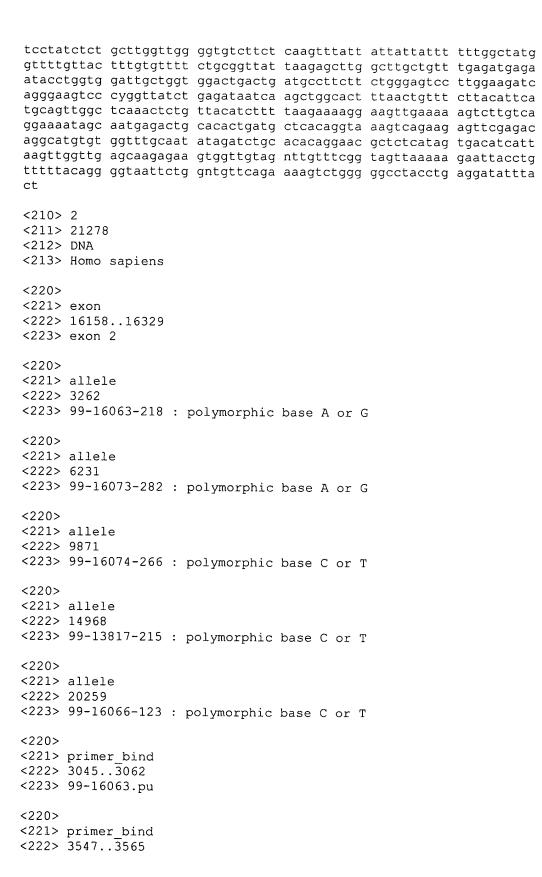


SEQUENCE LISTING

```
<110> BLUMENFELD, Marta
      BOUGUELERET, Lydie
      CHUMAKOV, Ilya
      COHEN, Daniel
      ESSIOUX, Laurent
<120> Genes, proteins and biallelic markers related to central...
<130> GENSET.045AUS
<141> 1999-10-12
<150> 60/106,457
<151> 1999-10-30
<150> 60/103,955
<151> 1998-10-12
<160> 71
<170> Patent.pm
<210> 1
<211> 5222
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> 1076..3075
<223> 5' regulatory region
<220>
<221> exon
<222> 3076..4643
<223> exon 1
<220>
<221> allele
<222> 4872
<223> 8-58-301 : polymorphic base C or T
<220>
<221> allele
<222> 3606
<223> insertion of AGAG in SEQID4
<220>
<221> primer_bind
```

```
<222> 4572..4587
<223> 8-58.pu
<220>
<221> primer bind
<222> 4990..5005
<223> 8-58.rp complement
<220>
<221> misc binding
<222> 4849..4895
<223> 8-58-301.probe
<220>
<221> primer bind
<222> 4853..4871
<223> 8-58-301.mis
<220>
<221> primer bind
<222> 4873..4891
<223> 8-58-301.mis complement
<220>
<221> misc feature
<222> 148,686,902,1258,1322,2440,2794,2852,3018..3023,3052..3053,3064
      3067,3428..3429,3489,3545,5131,5182
<223> n=a, g, c or t
caatatacag gtcacacaca acacagccca cttatctgtc agtagtccta ccagtgccct
                                                                     60
gggctccttc aaggtcccaa gtccttgaaa tttgctgtcc ttctgaattg tccttttctc
                                                                    120
ctccctagaa gacttctttc tgctcccntt aaatgtcccc aattcctatc tatccctagt
                                                                    180
ccctcttaaa atccaactgg agtttactgc tctactctga agtctcaaaa catgctaatc
                                                                    240
ttcacagatt accttgtatt gtaattgttg gcctagttag tattatctgc cccttgacta
                                                                    300
agcttcaaga ggccagactt ggtcttttat ctgtcttcca acagttcatt ctagaaccaa
                                                                    360
cacacaacaa acaaaagtca aattggttct caacagttta ctttcagtgg gcttgcgtcc
                                                                    420
480
ccagcttact taaaaataat taactatagt ctgttacata aagtaaaaat agattaaaca
                                                                    540
ctattatttt aatgcagtat ataagacttt tactaagtat cactgaacta ccttctgaga
                                                                    600
catgattttt actagatctg aatagtaata agataagtca cagagtcttc ttttgcttcg
                                                                    660
gttccttgtt tgataattat gaaccnttta aatttatatt gtttatttta tttcccctag
                                                                    720
ttttcgttct ctttctgatt taatcacaca tgcagagtaa gacattggat gctaataatc
                                                                    780
tcagccacca ttaaaatgtt tgtgtctgat taaaattaat qcattattqa aqqqaaatat
                                                                    840
attattatct aataagtgtg gatgtgtatt ctatcatact tcaatccttc gaaaatgctg
                                                                    900
gnactagtga atatcacaaa acaggaaatt agtaatacaa aataaaaatg ttaggaagat
                                                                    960
gctgtaaaat gaaaatactg attatatctg ggtgttgata tcaccatgac tttttaaatt
                                                                   1020
gctcttctga tttttacacc ttttttacaa tgcacaaact gtacttgtat aataatagtt
                                                                   1080
atgtcagcga gaatgtatgc actcgtattt agttaagaaa tattgtggca tgtacagtac
                                                                   1140
agtcctgatt aataaacaat ctggttaata aataataatt ttgtctgtat tatgaggaaa
                                                                   1200
aactgacatt actcagaaat tcaacacact gaaatcgaaa gtactagaat gatctacntt
                                                                   1260
acacaaatcg tcttttaaaa cactacatga cacttgagta aaccagccat caaacaggac
                                                                   1320
gncggctttt ggatgatcat tacctatagt tggtaagaac tgaaaatttt taggggtaaa
                                                                   1380
atttagtgta tacagtacct tattatgcag cagaagaggc tagtaggtcg aatcctgccc
                                                                   1440
```





4740

4800

4860

4920

4980

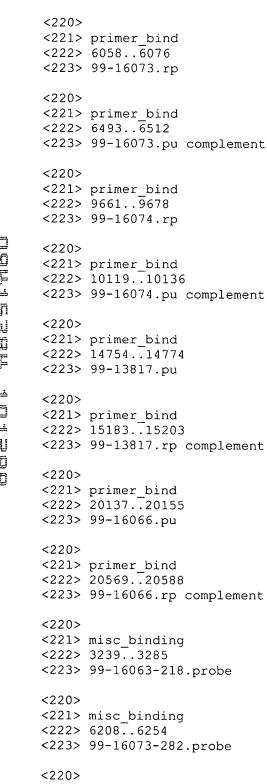
5040

5100

5160

5220

5222



<221> misc_binding

<223> 99-16063.rp complement

```
<222> 9848..9894
      <223> 99-16074-266.probe
      <220>
      <221> misc_binding
      <222> 14945..14991
      <223> 99-13817-215.probe
      <220>
      <221> misc_binding
      <222> 20236..20282
      <223> 99-16066-123.probe
      <220>
      <221> primer bind
      <222> 3243..3261
      <223> 99-16063-218.mis
      <220>
<221> primer_bind
      <222> 3263..3281
m
      <223> 99-16063-218.mis complement
Ш
      <220>
      <221> primer bind
      <222> 6212..6230
      <223> 99-16073-282.mis
      <220>
      <221> primer_bind
      <222> 6232..\overline{6}250
      <223> 99-16073-282.mis complement
      <220>
      <221> primer_bind
      <222> 9852..9870
      <223> 99-16074-266.mis
      <220>
      <221> primer_bind
      <222> 9872..9890
      <223> 99-16074-266.mis complement
      <220>
      <221> primer_bind
      <222> 14949...14967
      <223> 99-13817-215.mis
      <220>
     <221> primer_bind
     <222> 14969..14987
     <223> 99-13817-215.mis complement
     <220>
```

```
<221> primer bind
<222> 20240..20258
<223> 99-16066-123.mis
<220>
<221> primer_bind
<222> 20260..20278
<223> 99-16066-123.mis complement
<220>
<221> misc feature
<222> 285,982,2264,2273,2823,11227,11232,13663,13681,13842..13843
      16849..16851,17022,17091,18127,20930,20962,20994,21207
<223> n=a, q, c or t
<400> 2
gggcgcatca gttctgggag atgaactctg gcttcacttc agtcagggac ttaaactttc
                                                                     60
tgggcctcgt attttcaaac ccatggaata gaggaaaaat attactgtct acctcagagg
                                                                    120
tttgtgtaag aattaaatga gttactacat gtaagacctt tatctagttg ctggcatgca
                                                                    180
gtgagtgcta tgccagcatc agtagttatc ttctgctttt gtaaatatgt tgcattggtg
                                                                    240
agatttgaaa tcaccgtttc aaattcacta tgatcattta tattngctgt gcgttgatag
                                                                    300
tggttataaa aagtcagaaa cccgatgcta agtatgtttt acattatagt ctatgaaatt
                                                                    360
cgagcaacag gattttatgt tttgtcatgt actcattttg ttccacagtg aataagaatc
                                                                    420
tgtttcaaat gaacttttac atttatacca tgtgaaagaa acagattatt qqataatttt
                                                                    480
tcattgtaaa agaaacaatt tccatctaag catttacaca ttctaataaa tgtatagtta
                                                                    540
acgttatttt gtgcttacgc tgaaaaatgt aaaagatatt ttctggcttg acctgatttt
                                                                    600
660
tttctgaccg cctctttcca taacatacaa ttggcttccc tttctgtctg ccttgtccaa
                                                                    720
ggtcttcatc caatgcagcc tgctctctgg aggctctgct qtttaqqtaq tgqtqtqt
                                                                    780
ttaaacccca ggaacatacc aaggtcaata gttttccttc cagctcactg actcttaaga
                                                                    840
aaacgacctt tatgcagggg actaaattct gtaaagacta ttgaagccat agttctaaaa
                                                                    900
aaaagttaga aacgtccacg tgtgtggcaa gtagaaagtt tttgtttgtt ttaatgtgta
                                                                    960
aggaatagct caggtccagt tnccaatgta agggaaacaa gcagaaggtg tttcattcct
                                                                   1020
tgataagcta acgacatctg acctaatgta tggaaagtgc gtctgtgcgc atgcgtgtgt
                                                                   1080
gtgtgcgtgc gtgtgtgtgc atatggcgag agtatggcaa tcagaattca ggggatgtga
                                                                   1140
tgctcctttg gtttgtgtgt tcagtatctc tatggaaaga gcaggttgat tgctaaqttt
                                                                   1200
actttgttat ttgagctttc agagtgagca tttaaagatc tattccatac acactgttga
                                                                   1260
ttgtctgggt cccaccccat ccccatcccc atcccccacc atttcgttgg ttgaaattta
                                                                   1320
agtttaaatg tcacctgagg ccaggttctg tggttcacgc ctgtaatgcc agtactttgg
                                                                   1380
gaagctgatg caggaggatt acctgaggcc aggaattcaa gaccaccctg ggcaacacag
                                                                   1440
tgagatccta tctctattaa aaatttaaaa attagctagg catggtggca tgcacctgtt
                                                                   1500
gtcccagtta ctcaggaggc tgaggcagaa agatcttgag ccagggaggc tgaggctaca
                                                                   1560
gtgagaataa aaacaataac aaaaacaaaa aataaaaaat aaggccaggc acagtaagta
                                                                   1620
gctcacgcct gtaatcctag tactttggga ggccqaqqcq qqtqqatcac ctqaqqtcaq
                                                                   1680
gagtccaaga ccagcctggc caacatggtg aaaccccgtc tctgcttaaa atacaaaaat
                                                                   1740
tagccaggtg tggtggtgcg tgcctgtaat cccagctact tgggaggctg aggcaggaga
                                                                   1800
atcacttgag ccctggaggc ggaggttgca gtgagccgag attgtgccac tgcactccag
                                                                   1860
cctgggtgac aagagtgaga ctccatctca aaaaacaaaa caaaataaaa taaaataaaa
                                                                   1920
taaaaataaa tgttacctga atattcccac ttaggtatga aattgatgtg cttgctttgt
                                                                   1980
tttatttcca tttgttcttt ctgttgttat tgttgttttt cagtgaggaa tcattaaaaa
                                                                   2040
tatcaaatta ttagctgtgg agactctcat ttacctaaga ataatctaat aactaatttc
                                                                   2100
ttttctaatt tttaaaccaa agttcagtgt gagatttgtg tattaaaaag tgcaatttgt
                                                                   2160
atcacaatca gtactaagcc ataaatcttt atcattgtta ttgatttaaa attatctttg
                                                                   2220
agccttgata tatttgcaat tgaatgaaaa taatttgtta aggngcttat ttnaggatcc
                                                                   2280
```



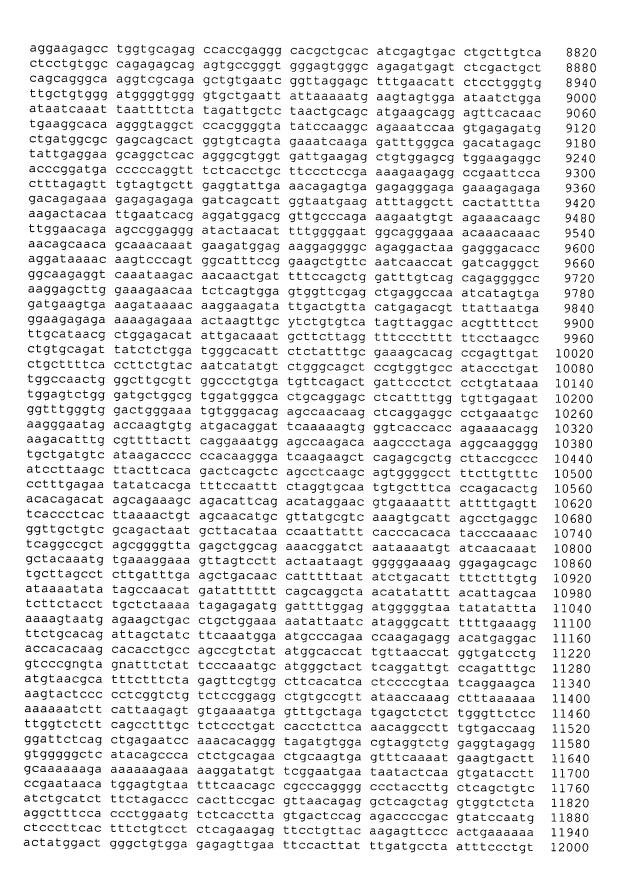


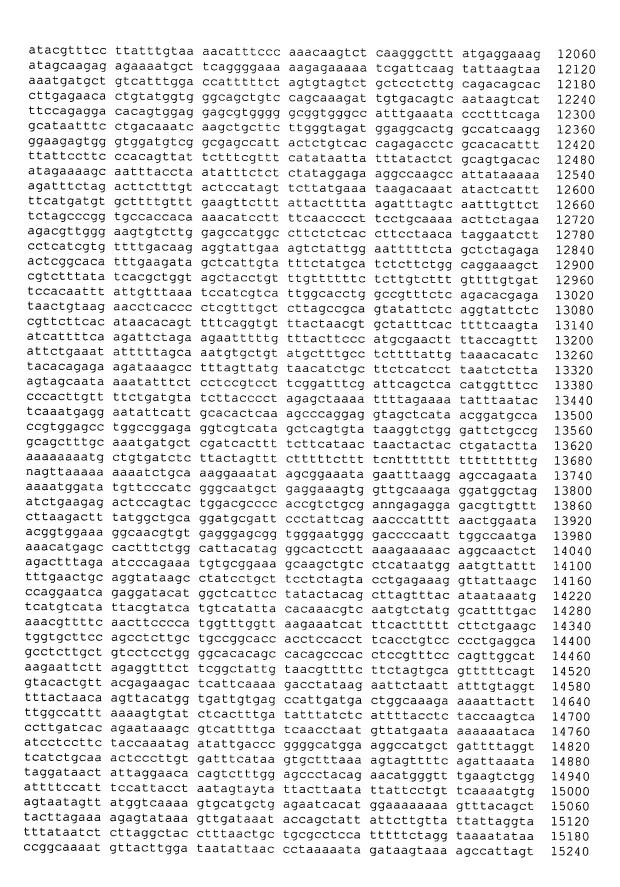
atggcagtca gcacggcaaa acacatgcta ggaaagtaaa tggttagttt taaattttaa 2340 gacattgcta cggacataaa atttattagc ttttttaacc agttaaccaa ttttttacct 2400 tttttaccaa ttgccaattt ggggctgaaa atataaatat atgtaatttg tattatattt 2460 atattatata taatttgtat tatatttata ttatatacat aattcaaaaa tqqaqqtatc 2520 atttgtatat acaatgacat atcttattga tgaatgtaaa catgcattct caagacagaa 2580 aattattaaa attaaatgac cagattttat tacatgtttt gcatcatgat atatgtgctt 2640 tattaatttg atttttgcat atttcagtat tattctacat ttttttgccag gctttactct 2700 gagcataaaa ctgggtgctg ccatttaatt ccaaatatat aatgatcaac tgtctgctcc 2760 aaagctgcac cttccaatgt taccaccaat tttattatta aggattgtta attccattga 2820 ttnatagttt tcaataataa tgaataatat tattcattaa tcaataatca actgataatc 2880 aataattaat taattaatca attaaacaat aatgttatga ttattcataa tccattattg 2940 tatactaaaa acatatttct gatggtatgt ttagctcagg ataaagaaga catgggctaa 3000 gataggccct ctgtttctgc caaagtagca gagaaaagtc ttctttccta gagttttgtg 3060 ccctaagcac atttgcaggg atgggttcaa agacttgcag agtgtgtact tatattctta 3120 3180 agaaaatgca ctgtattata ctaacttcac aggggttaga ccgcagcaag tcaatggaac 3240 acagggagca cccaacgaga arctgtcctt gaaaatgttt tagggaggaa ttctaaagtc 3300 ttgcctttca acttaactgt gtatcttcat ttgaattatg gaaccattgt tcatgagtga 3360 aatctatttt tatgttctga atcttcccat ttaaagacat atgttttctc agcttttaga 3420 caataacatg atctaaaatg tcaaaataat acattctgct gaatctcagt atttactgag 3480 gaaatacaga aacatggcaa aagttaggaa gaaaaatgtt tcctttttct gctgtccaca 3540 aaagttggtg gaacatttat gactgcaaaa aaaaatgagt tattaacact gttttgattg 3600 aattcaacag atgctctgac tttcttatat atatgtgtgt gtatgtatac atatattt 3660 aacaactata aatatttata tgtaacaaca ctttcacaag atgccaataa atttagtatg 3720 agcatttaca tatctaaatc tttctttggc tctagaaact gatgctatgc caaccagaac 3780 aaagaaatag tagctttcac tactgtcact gctaataaaa ttcagcatca ttacagctta 3840 tacaaaaattg tacttattag cccaatttct attgcttgcc ttaaaaaaat gtatcggaga 3900 aaatggatgc cattagttta gtttgtagat taaactgata ataataatgg caaaaccttg 3960 atcaagtaaa taaatcccac taaatagaac aggaatgtgt tgctaagacc aaggccattt 4020 ggattcaccc tgagcttgtg tttagaggca ttctttcatt cattcattca ttcattcagg 4080 cactaatggc tgagtcatgc cagaggctag gcccctcctg cggagagttc actgctgcct 4140 aaagacagac caggttcttg cccttatgtt aatattcggg tcttgcctag tcagttatat 4200 aaagttcatt aaaatttgct tacataattc aagcaagtgc tcagttttaa atgtctgtgc 4260 cacacaattt gacaatttac ataatttatt tcagtgctaa agaagttgac aatcgctgtt 4320 taacttcagt cgggcttctc ctaagcgggg cgaacacaca caggacatgc acaagataat 4380 ctgttaaagt atgacaagaa aataataaat tctacttgca gttataattt tatcctaacc 4440 tttgcaatca cttgtgctct gtgtgtgttg tataatgtac acattatata acaqttcatg 4500 tgtatataaa tgaatattta tgtaagggaa atatgctaaa aatcttaaat gggtggaata 4560 actactaaaa atgtttagca agtacaatta aaacatatac aagttagtgt cacttctgta 4620 taatttaata gttaaatagt tgtttctatg gggaatagta agccacagag agaaagaagg 4680 tettaagace gggaceagaa aaataagett ettaattgga ggeggeggtt aettetagtg 4740 ataggcgatt gtttttcca gatataattc tggctccttc ctaattaaga ttagtgttac 4800 tcatgtcacc aatttgcagg gtttcaagct tgtgttaata agatcacaaa cattgtgagc 4860 ctttgccatc tttaaaatct ttcactctaa agcaaaacgc acataacagt tgccgaagtg 4920 ccttctttga gcttctttga aaacagccgc attttctttg agctatggag agcagcatag 4980 aacatgtttg aatccccaaa gcttaacccc agaaaagata tttcttgcaa gccacaaaca 5040 attctacact tacagtatca tgagttaata tttacctctg cataactggt atagaataaa 5100 acacagtttg atgtggggtg gcggggtgta gggtaaaatg acctaaaaqc ccaccctqtc 5160 tctgttacat tttgctgtgg gactttgtcc atattgctta cattctctga gtctcggttt 5220 tcatattcat aaattagaat aatatcactc acatcacaga taccgtgcat tgaaaacacc 5280 tggcaaaata actgatagta aatagttctt ttttattttt ttacttttta ttttttgaga 5340 cagagtetea etetgteace caggetggag tgeaatggtg cagtettgge teeetgeagt 5400 ttctgactcc tgagttcaag tgattctcct gcctcagcct cccaagtagc tggaattaca 5460 ggcatgcacc gccacgccca gctaattttt gtaatttttt agtagagaca gagttttgcc 5520

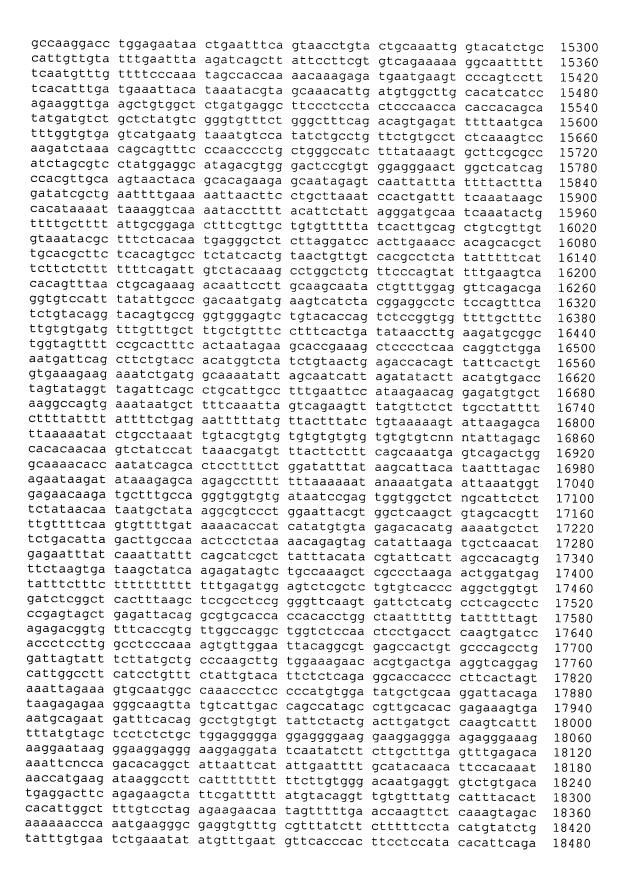


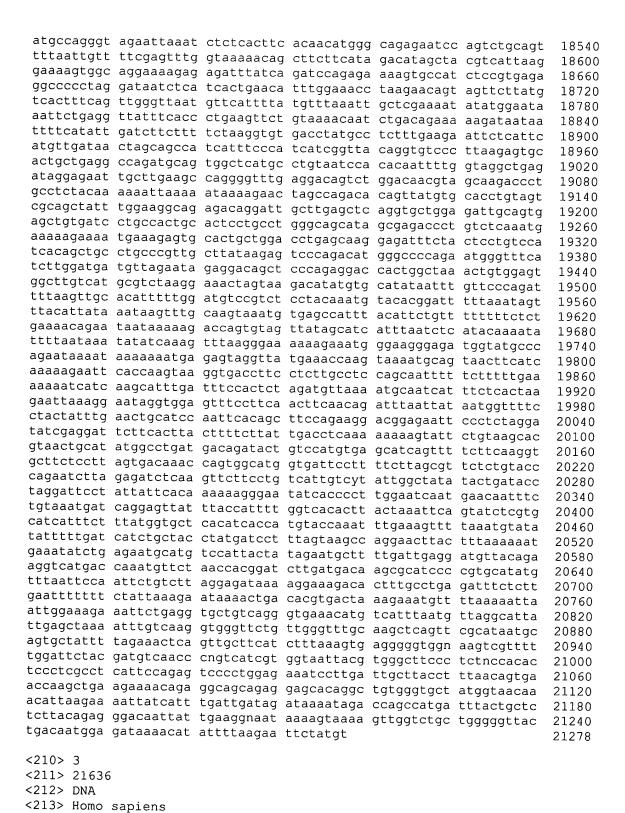


aggttggcca ggctggtctt gaactcctga gctcaggtga tccacccacc tcggcctccc 5580 aaagtactag gettacagge atgagecace atgeetggee ttgaattttt taaattteaa 5640 tagctttagt ggtgcaagtg gtttttggtc acatggatga gttgtataat ggtgaagtct 5700 gtgattttag tgcaccggtc acctgagtag tgtacattgt accctacagg tagttttcta 5760 teteteacee ectteacaaa etececeatt etgagtetee aetgtetgtt gtateaceet 5820 gtgtaccttt gcatacccca taggttagct cccacttata aatgagaata tgcagtattt 5880 ggtttttcat tcctgagtta cttcaatgca aacttgctgc aaaagaaatt ttgttctttt 5940 6000 ttcccttgct gtcttcctcc catgctctgc ccttcctccc tctctttctc atatcagaag 6060 cacactaacc tccaccccac tcctctct gacatgccca tctcactccc tcacatgccg 6120 tectettete ttetecagaa acceteagee eccattgaet tetgeteett ecacaeggag 6180 gtgcactgac gggagcattc acagtgggat tcatcttacc ctggctcctc raattttaca 6240 gcactatttc tgatgccagg gaaaatgtct tatgatgaaa tttttaacct gccctacacc 6300 tetgeteece aaggeaceta tetaetgtea gaeteeaaag ttggetetea gtgaaaaatg 6360 tgggattaac agacagattg gatggtacct aggcagccag tcctctgttt catggtaaca 6420 ggctacaagt ctgtatttca attagaatac tgaagtgttc actctgactt tttaaatatg 6480 taaattctaa tagaatcact ggcatttttc tgacagtaat gcagaccagg tttaaatagg 6540 caacatgttt aaagaaccac gtgacacaat tctactccat tcccagggtg ccggttggct 6600 cttaatttac aaaaggggtt tgcattcata cctttcccag cttctcacta gcaggacaca 6660 ttaagataat gcagaaaaca acaaagatag tgtggtgtta gctgctgttt cttgaggagc 6720 aaatgcaatg acaaaatcag aaatgctggc atctgtggga aaaacaaaga atgcaaagag 6780 cctgctccac tgaggaacag cagattttct ttgagcaaaa gcagattatt cttgtttcaa 6840 ttataatttt tgtgactgtg acattggtaa agtagggtgg caatcatgca ctcttggtac 6900 acaagtgacc tgacaatatg gccaagaagt aaaactttta gatacatacc aaaagcatta 6960 tatattctag ttctataatg tgcacagcta tgtaaaagta tcagattctt agataactaa 7020 ctatagttta ttatgtcaca ttttccatcc tttactttgg attatatgac aaagttatgc 7080 attaactttc ttcagactct tttttttgcc catttctact aagtgaggct gttacagtaa 7140 gttcctttct ctttggtttt atgaagaaac gaatacatct gcattttata gtgaatttac 7200 tgattactta ttatagctct actcagtgcg ttggtattat tggtattttc taaatagatt 7260 gtagattatg gaagataaca ttcatttatt aatcatcaag tgctccatga aatcttatat 7320 gagttctgga aaagaaggaa agcctaacac attctctctt ctcctatcct tggtgaatat 7380 attatccctt tatatcctta gattagtttc agatttctta tttaagactc aattccattt 7440 ccctttcttt agaagactga tttcccactt tttaaaccct tttcaacatt ctcaactcac 7500 tcaagatgtg gagcccaaaa ttgaatttat attgaagtgt tcacttgctg cctttttaga 7560 gatataaatt ttaataggat cactggcact tttgtgacag tccatacagg ccaggttaaa 7620 taagtggtat gtttagagga catgtaatac tetgcaette taaggeattg ettgttgget 7680 gagagtccac aaaaggtgct ttcatttact acgggcccat tccgactcta gataaggcta 7740 ttgagttata tecatecaeg gecaaeaete agaagtagte ttatggagtg aetttettet 7800 tttttgatgc ccacagcaga gctcccacca tcttcgtcac taacatgata atccggtgaa 7860 ggcagcttgt taaaagagat cttccttcac accaaaggaa acgcttcctt atttactqtq 7920 aagttaattc actggttatt atcatcgtta atattcattt ctggatatat aatagaactg 7980 ttctggaaag gtatgaatta gtataacctc tctgctcaac actttttgcc acaatcccag 8040 ccaaagtagt gtgtccgttt aaataacagc tataaaatat actttgccct ctgaagctct 8100 gtggccttcc atgagctcag aacaataaac cacaccattc cccatctttg tgtgacaatc 8160 tttctcagct tgcaacccaa tcccatattg tgctttagga cctgaaccat ggggtctttt 8220 tggaaggaga ggttcacttc atgaagtgga cttttaaaaa tttattctt tagtttttgt 8280 ttgtttgttt gttttgtttg tttttgtttt cggccttcaa taacggccac tgggttggtc 8340 atatgaaaag ggattcttag gatttaatat gttgggggct catcagggtt cctggtttga 8400 gtctctgctc tgggcaggat ctcagggcca ggtggactgg ctctcacqct aaccaatcqc 8460 ggggcatcat tatgcaagcc caaaggccat gttactagtt tccagtgggc attccactga 8520 eggtaaattt etgeeagata eateeaggge aetggggata eaagaaggea eaaagettgg 8580 cetteattee aagageatag ggtetgggaa ggaggeagae gagaagaeag eateaeatgt 8640 actgaactgc cgaagaagag ctcattgcag ggtctagagg gaatgcagga aaggcttcca 8700 agtgaggtca cacaggatgg acaggaatta ataggctgag ggattcgtga gggaggtgga 8760





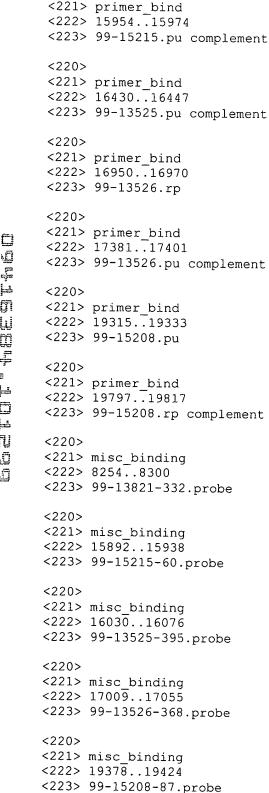




<220>

```
<221> exon
      <222> 5537..9359
      <223> exon 3
      <220>
      <221> misc_feature
      <222> 16330..18329
      <223> 3' regulatory region
      <220>
      <221> allele
      <222> 8277
      <223> 99-13821-332 : polymorphic base C or T
      <220>
      <221> allele
      <222> 15915
<223> 99-15215-60 : polymorphic base A or G
<220>
      <221> allele
m
      <222> 16053
<223> 99-13525-395 : polymorphic base A or G
      <220>
      <221> allele
      <222> 17032
1
      <223> 99-13526-368 : polymorphic base A or G
⊨≟
      <220>
N
      <221> allele
      <222> 19401
Ø
      <223> 99-15208-87 : polymorphic base A or G
Q
      <220>
      <221> primer_bind
      <222> 7946..7965
      <223> 99-13821.pu
      <220>
      <221> primer_bind
      <222> 8454..8472
     <223> 99-13821.rp complement
     <220>
     <221> primer_bind
     <222> 15475..15495
     <223> 99-15215.rp
     <220>
     <221> primer_bind
     <222> 15943...15960
     <223> 99-13525.rp
```

<220>



```
<222> 8278..8296
      <223> 99-13821-332.mis complement
      <220>
      <221> primer bind
      <222> 15896..15914
      <223> 99-15215-60.mis
      <220>
      <221> primer_bind
一一
      <222> 15916..15934
      <223> 99-15215-60.mis complement
m
      <220>
<221> primer bind
      <222> 16034..16052
      <223> 99-13525-395.mis
a
      <220>
Ի
      <221> primer_bind
<222> 16054..16072
ᆣ
      <223> 99-13525-395.mis complement
N
Ū
     <220>
      <221> primer_bind
Ū
      <222> 17013..17031
      <223> 99-13526-368.mis
     <220>
      <221> primer bind
      <222> 17033..17051
     <223> 99-13526-368.mis complement
     <220>
     <221> primer_bind
     <222> 19382..19400
     <223> 99-15208-87.mis
     <220>
```

<221> primer_bind <222> 19402..19420

<221> misc_feature

<220>

<223> 99-15208-87.mis complement

<220>

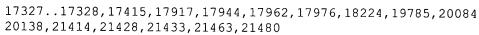
<220>

<221> primer_bind <222> 8258..8276

<221> primer bind

<223> 99-13821-332.mis

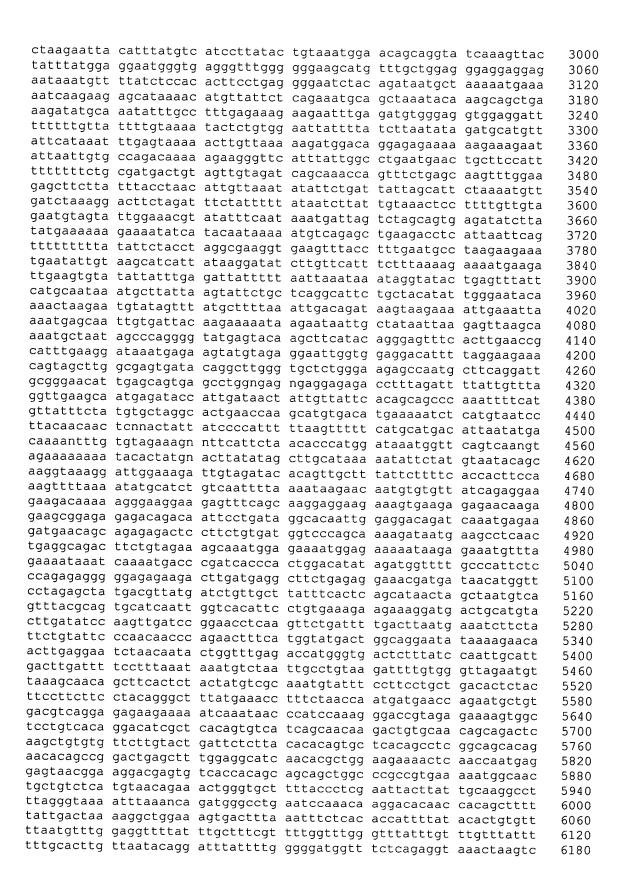
 $<222>\ 110,\overline{4}287,4291,4453...4454,4506,4520...4521,4558,4580,5958,17188$



<223> n=a, g, c or t

<400> 3

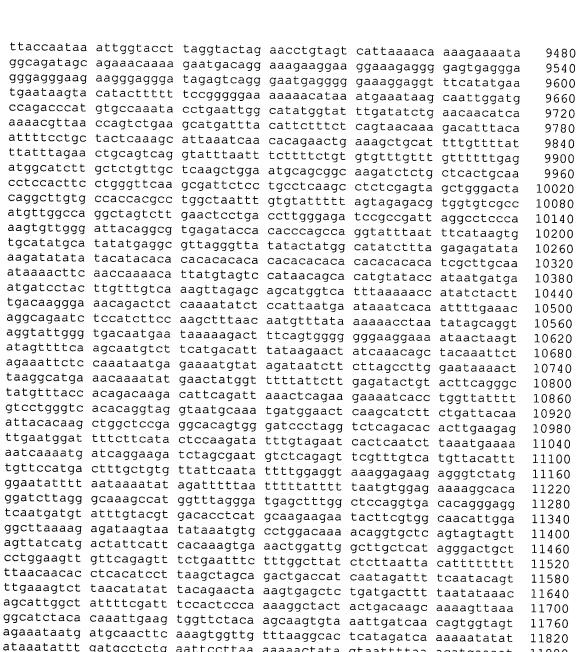
aaaaaaaact ctgtgaggtt ggtactgtaa tgtcaaacag caggtaatga ttcactagag 60 atttcaaata gaacagctta taagcaaaat atatcagaac aacatttcan tacccggaaa 120 atttgctttt gtttttcct aatgctaatt agtgcaattg tacttgtttg tgcaaggagg 180 ttcttgtttg agaaattatt tttgtttgga aatatgtttt tataatttct tcaatgctga 240 ggaaaaggtc atttctgtag gtaggctttc ctaggaaatt tgtcccagca ggttgtgtgc 300 aggtggtttt tgcgggagaa ctctcaggca tgctacctat gagggggtga agaaggagaa 360 420 gtcaagtgga cccagccttg ggacctgtac atcgttagga ggctgcctag ggcctgagaa 480 ataggtcaag actgctaact tcaaccaagg gcagtgtccg ggaggaatgc agggtgaaac 540 catctgtagg tagcactgcc aactggagga gtctgtgccc ccaggtccca agggtgtcat 600 tgatctgctt atgtgaccta ctgactctat ataaggcaaa gaaataatgg ccttagaagg 660 ctgcatgtcc ttgcttattc agaaatgcat ttgacagaga aaaaggcatc tgtcagaatt 720 caagtcagag aagaatgatg tagaaaaaag agctggaaaa ttttctaggc agtcccaacc 780 aaaaaagacat tgttttatgg cataaatcca aaagaatcaa agaataggta aggagacaat 840 aatcaaaacc tcttagaaat tagaaagttg atgcatgcta acagatttgc tqaaacttag 900 aaaaagtaat atgaagtagc agagaggaaa cagagtcagt tcgatttaca tccagagtca 960 ggtattgaca acacgagatg caatggaaga aagagcaaaa taaggaagac tgggtggaag 1020 tcagttttaa gaggcagtca ggtccccaaa tcccattccc tactcccatc ccagagactg 1080 ggggcatatt tcttcctgga gggaactttt tctttttgag atgataaaga aaaggttggg 1140 ccgggcgcga tggctcacgc ctgtaatccc agcactttgg gaggccgagg cgggcagatc 1200 acgaggtcag gagatcgaga ccatcctggc taacacggtg aaaccccgtc tctactaaaa 1260 atataaaaaa ttagccgggc gtggcggcgg gcgcctgtag tcccagctac tcgggaggct 1320 gaggcaggag aatggcgtga acccgggagg cggagcttgc agtgagccga gatcgcgcg 1380 ctgcactcca gcctgggcga cagagcgaga ctccgtctca aaaaaaaaag aaaaagaaaa 1440 agaaaagatt gttgatctga gcaacagtag gaggagttca gggcggggga ccactctgac 1500 caccaggatt ccatgcacaa ctccaccctg aagctgagac tcccccagcc acatcctgct 1560 teteceteet geteeetgge ettetgette etggatggat attggataga ettttetggg 1620 aagtctgatc aacctaaaga cagaatttca aaagttcctc catcctggcc gggcacggtg 1680 gctcaagcct gtattcccag cactctggga ggccgaggcg ggtggatcac gaggtcaaga 1740 gatcaagacc atcctggcca acatggtgaa accctgtctc tactaaaaat acaaaaaatt 1800 agccaggtgt gttggcggga gcctgtagtt ccagctactg gggaggctga ggcaggtgaa 1860 tcacttgaac ccgggaggca gagcttgcag tgagctgaga ttgcaccacc gagctccagc 1920 ctggcgacag agcgagactc tgtctcaaaa ataaataaat aaataaaaat aaaaataaaa 1980 gtttctccat tctaacagtt caggcatgtc atctcacgtg cagctcacag gccgcagccc 2040 tccttaggtg cccagacttc ctgtgcttgt tattttgtgc agacaactag gaatgcatgg 2100 tgtttccatg agaagaacaa gggacaaaca aacacaaaag ggtggcaaca cagagaaaat 2160 ggagggaatt tattcaggga gggaatgaaa cacaacaaat aagcgtaccc caaatatctc 2220 agagtgagaa aaacataaac atggtaacca taaaataaga acagttgctt cgaaaagcaa 2280 cattcggata acacaaagga gctcttggaa cttgatccta gggaaaaaag caaaattcag 2340 tgaaattatg tcacgtgaaa actcaccaaa ttccctgaaa gcagagttaa gggatgaagt 2400 aatatgaaat attaacagca gaggtaagaa aattaaagga caagttctga aggtccaaca 2460 gccaaacata ggaccagtgc ttgaataaat ctgaagaacg atcatttcta aattagaatt 2520 ccatacccag ccaaacaatc aattaaatat gaggtagaat aaacacactt taaggcatat 2580 ggcatcttgg aaattcgccc tcatgagctt ttcctactga agacactaga ggttgtaaac 2640 cctcaaaagg agaaagaagg ggatgaggca caggatacag ggaacagggg atccaqcact 2700 gcagtgcacg gagcagtgga gactcatcag gcatgacggc aaccagtcca gatggtggga 2760 ggctgcagga gatatttctt cagaaaagaa ctgatagaac acttcatgca gctgaaagaa 2820 ctggcaatgc atttgggttg aatgagttag aagagggaaa ataacaagac aagtattatc 2880 tgtagaaaaa aaaacgaagt ggtataggga tagaaaattg gactttacta catggttgag 2940







ttttcactg	t ctctatctct	: ctatatattt	: ctagtcattg	ı tgtgtgttca	tcagatagtt	6240
ctgtcttta	t gtcctgtcag	, cttctattag	, aggaatgatt	gctatgacct	catggtatag	6300
caaaaaaca	a caacaaaaa	ı agaataaaaa	ı ataaaaaaga	caaaaaaaaq	aaaacaacaa	6360
aaaataaaa	a taaaaaaaat	ccctaagtct	cccttctacc	cacagaacca	acaacaccct	6420
tcccagcct	t tootttooct	cgccctcttc	: tcgtccccta	agcaaacaac	atccacttac	6480
ttctgtctg	t gtaaccacag	r tgaatgggtg	r tgcacgcttq	gtagacctct	gagecetat	6540
tgcacaaac	c agaaacagag	cagagccaag	ggggcctgac	: aagagttcct	ttttagctga	6600
acaaacaag	t gctctccata	. ataggtggaa	tcagacagtt	aacacatttt	tatottoaaa	6660
acaaaataa	a aggaaaaaat	taaaaaaaac	: tatcatgaac	tgtattgctc	cagttcccat	6720
ccccaagtg	g cccagccctt	tcttgctggt	ccagctggac	aggagcagct	atctagaatc	6780
aggatgcgg	g gagtgaggaa	gtttttcctt	tgacaatgaa	agtagacttt	cattotoatt	6840
tttgttctg	t tgcagtaata	taggagcaca	ttttggccat	tgtaattaca	gggaacaaag	6900
ggattgcgg.	a cacatatctg	gacttctttt	cctcccttat	tgttgtggaa	gagacactag	6960
aaatgctca	a acacctgcaa	tatacagaat	atacacaatt	ttattccagt	atttccctaa	7020
catatggtt	t aaaattattc	caggtataca	gtgtatgcaa	ttctgcatta	tcacagagga	7080
acaacttct	t ttttaaaaaa	taaataggtc	agccattttt	attaacgtgc	aaaaacttta	7140
tcactctaa	c atgctctagg	tagttgagga	aaagaggtct	gatcactgtt	tatatttat	7200
tttctttgt	g ggaacatttc	acctgctgag	tgtacatgaa	tttactttct	ataaaaaacct	7260
tttatgagt [.]	t tacagtagaa	tcagtggaag	gaagagttaa	taagggctgt	ttttaaaaaa	7320
acaaacaaa	c aaacaaaaca	aataattaaa	aaaaaatttt	acattccttc	ctattctcta	7380
actacactt	g ggaagtgcac	ttcagataag	tttgcagtgt	gactgagaga	taaaaaaat	7440
ccatagaaa	ggtcctctta	gtgaacaaaa	tttagttatt	aactttatag	ctatgaaatt	7500
teceeggge	tttgtttttg	ttcaaacaga	ctttaacctc	tacatcatac	ttaaccetee	7560
gacatgcgta	a cagtatgcat	attttgttt	gaaaaaaaat	atttcattcc	actotattaa	
gaatattcaa	a aaataataaa	ggtattgctt	aataaaatto	ctagaattat	ttagaagtag	7620
atqcacaata	a ttttactaga	ttctttattt	taatagtgt	ttattazasa	tagcagtac	7680
aaaatggtct	gcgcaaatac	aaaaaaaaa	aaaacaccaa	aaatacaaaa	tydadaictt	7740
tttgttcct	tttaaaaatt	tttttttcta	caaatacaa	tacattcaca	tataaaaaa	7800
tacatacaca	taggcatcat	attttagtgt	atogaaacat	ggtggtttta	tgtggacaga	7860
agctgaggaa	gtgaatggct	caaaaaaaa	acggaaacac	ggtgettte	tggggacaca	7920
aacacttgta	gaaatgagca	ctttagtatc	ccaeccaatc	catggaattg	cigacigiga	7980
acactagaaa	gacgagaagc	agaggaattg	aggetgeaag	tasttastas	gatttcaatg	8040
acagggtgat	tttattcaga	tracctctaa	aggetgeaag	agazaga	ggaagagete	8100
ctttatctac	ctctgctgtg	gtaaggccag	togatotogo	ayyaayyayt	ctgttctgac	8160
ttcctcatgo	tgcttctccc	cataccataa	gasttagat	tattaaaaaa	ctcttcagcg	8220
gcaagtgcaa	attccactca	cttttattag	tataaaaata	tettgeagga	aaatgayaat	8280
tagcaaaato	acceaeacca	gtattcctta	nacyacaata	aattattett	tctaaacact	8340
ttgaactgac	acgaaaagca taaagttctt	tacaaacaca	adatataggt	tttcattttc	tcaatatatt	8400
tttagcgttt	atttaaaaat	aataccaaca	agaggatcag	aaggttccac	ttacccgtgt	8460
ttacataaga	gtttgaagat	acceccace	aaaaygtgga	aaactaaaac	caccattgct	8520
gagaaacact	ggtacacact	ttataaatat	CCCadadada	aacacagctt	tgaaagttgg	8580
taaggaggga	gcagtcttca aggggaatga	atactttata	adayiyalad	tttaggtgag	gaaggaacgg	8640
tctaattctc	aggggaatga	ttetttete	tattastata	cagecageat	tggaatagtc	8700
tttcagtgc	gcggaatatt	tagtgtaa	gttcatcta	ttcttattac	aggagaatga	8760
ggacatgtto	tagtatttat	cattetteta	gigiaaccac	tttcccgcaa	tatttccact	8820
actectttee	aagcaaagca	gerrerrer	gggtcaggca	gaggatccaa	ataggcaacg	8880
tctttataat	caageteaga	aacaaccact	gatgatgcag	atttcactac	atccgttgat	8940
tttgaacaaa	agttttcctg	gcacacactt	gagcatgctt	aaggttgtgc	atatgttcat	9000
atagaggata	taagttcagc	ceetgeteet	ttgtaatcat	catcttgttt	agtttttcct	9060
grayayyary	tctaccgtat	actggctaat	gaactacaat	attaaaataa	ttatttctgc	9120
tatttattat	aaaaccaaga	aaacaatatt	tgtttcataa	atgtatggca	tagatttccc	9180
agt >> ~ ct = -	tctttttatc	rgcaatagag	aaatttcagt	ttctggaata	tgttggtact	9240
gctaagctaa	atggcaaata	acatagagtt	caatgtcatt	ctttattgca	tttacatttg	9300
gryaracytt	atattgaact	tttcttgcat	taaaaagata	gtatttttcc	cctctctcac	9360
aayycaytaa	gtgaaagttg	ttcttttaaa	tggcatcatt	acaaaaagta	atttggtttc	9420



ataaatattt gatgcctctg aattccttaa aaaaactata gtaattttaa acatgaaaat 11880 gtttaaattg gaacatttat tatgaaaatt aattaaactg aattaatata tattttaatt 11940 aacactataa ttattataga atgaaataac agtgaaaaag ttagaaaacc tgttcagttt 12000 ttgataaagt atatggaaat agattaagca gggctcttga gtaaattatt tcattgattt 12060 gtatctattt aggtaggtga gagttgcata gaatcagtgg cgagaaggat cagtgaaata 12120 aatttacaac ttcagtgaag accaggtaat gagcattgat ctatgaatac agactttctt cagtgactat ttcccctaaa actcccagct caccgcagca ctgctgcttc aagattccag 12240 ttattggcgt gctagccaag tagaaggtca tttttaccga aacgcaaatt tcaacttatg 12300 gccattttct actgaccatt tcactgcaaa atggataaaa caataggaaa tataaactac 12360 agcaaaagtt aaagctctat atggacgtta aaagaaaacc tcttgctttc tttttcttcc tttttctctt gttttctttt tctacctttt tctgcatatg gcatttattt tcatatcatg 12480 ataagcattc tcatgtttgc acttaatttt ccttatttct atttttctta gatttgttga 12540 tgttgtgtgg catattaaaa gtaggaagga ctgtataaac tcatgtatta tttttaaatt 12600 gtttttctga atgtactcaa gtacttttt atgtgaatat gaacacggaa taaaagtact 12660



aaaaccagtt aattatgaga gactaagaat gtggaaagga acattccgga gtatattagt 12720 agctaaagaa atgccttgga tcttaaattt caacttctaa aattattaaa atacaaatat 12780 taaaaagata tagaggccac tgctaaagtt tgtttacgcc ttatttcttc ggttgcctag 12840 attattcgtt ttaggtgaat ttgctttttc tgtcaccgtg gttatgttaa cccaactgat tcatccaaaa cggaattggc taaactcctt tatgcacata tttatattag ggctgattat 12960 tttgctcatt gagtttcaaa aaattatcta gaggactgcg aagttttcat tgctttactt 13020 aatttctttt tcgcttgtgt gatttttatt tcttactctt gactatatct ctatacggtt 13080 tgaagtaaca gggcagaaag ttaagctaat aatttctgct gacttgataa ttcatcagtt 13140 tgacaaaaag accagtgggc caccattagg aactgaaggc aactgaaatg acatggctta 13200 ctcttgctgg aatcttagct tgaaagttga acatctactc cacaatgcca cgggatccgt 13260 tcatgatatc atttgctgaa gcacttgcca agaatttcat tatttagaaa aagcagaatt aacatttcta tctattattt ttctttctt tctttaagct gaaaggcatt cagaggctgt 13380 cagtaaataa aatctatcag gaacaaaatc attttcaaaa tttttaaaca tgatttattc 13440 ttgatgcaaa tcttattgtt cttacaattc tatgcaagtc aagttgagct aaagcatttt 13500 tatttaattc aattttcttt cttctgaaat gtgatacagt aaaatcaggt tttgaatatg 13560 tacaaacaca tcaggatgat tattatactt acataagtaa aatatatgta aaatattatc 13620 atatcttcac catatttaga gtctaatatt tgttagattt tttgtgtgtg catttttcct 13680 ttccccttca gaaaacgtag ttttttattt aattcatgtg gtttttctat agctttctgg .13740 gtctcatcaa atttcacatg aaactatgca gagtcatatt tatgttaagg aagacatgca 13800 attaaaaccc tacccatttg ttttgttagt cccaacacgt atcaaacaaa aaacttatgt 13860 tgttagatgt agatttgatt atctcccttt atttatttgg gtctctctct ctctctct 13920 ccccgcccca ccgtctgtct cactctcact ctctctctt cactttctct ctgatctctt 13980 ttcctttcaa ctcagcttgc aggaatggat ctcatttgaa atgagctgca ctggtgtcag catggttgtt taaccatgta aactatctca tgatgcttag aattgaagag aagcttaaac 14100 gcttgtgtca gaaaaatatc ttacaataca caaaacacat taaatatatt attagaaaaa 14160 gtttagttgt gcaacaaaga cctttaattt caactaaaaa ctgatgcagt aatgtattac 14220 aatgacccaa aaagaaatat tottttcaaa atgacttcgg gaagctatca gagtttgtgg 14280 gaaaaatgct aaggtaatac tttcctaaga acaaatgtaa agctgcaaac aatgactttg 14340 aatttttcaa gattagcagt ttggatttac atgtatttta acaaaatata aaaaatttac tttcaaatct tggctttaaa aaaaaatcgt gtggctcatg cctgtaatcc cagcactttg 14460 ggaggctgag gagggcggat cccaaggtca ggagactgag accatcctgg ctaacacggt 14520 gaaaccccgt ctttactaaa aatacaaaaa attagctggg cgtggtggcg ggcgcctgta 14580 gtcccagcta ctcgggaggc tgaggcagga gaatggcgtg agcctgggag gcagagcttg 14640 cagtgageeg agategegee actgeactee ageetgggea acagagegag actaegtete 14700 aaaaaaaaaa aaaaaaaatc atgtttcaac tgaaatggat atttattatc tgcaaaagtc 14760 cattttgttg catatattgt ttaaaataat tcttaagaaa ttgttacctt tctatttcta 14820 tggagaacta acaccttaat acaagtctag cccccttttc caagacacaa cgtcccccac 14880 aatgccactg agtttgtgtt tacctgaata ctttcaaacg acagctgatc cttgaataac 14940 atgggtttga actgcatgag ttctcttcta tgaagatttt ctttacctct gccatccctg 15000 agacagcaaa accaaccct cctcatcttc ctccttcaaa gcctactcaa catttgaaga 15060 tgatgaagac gaaggccttc atgatgactt catatgactt tcccttaata aatagtaaat 15120 atacttttcc ttccttatgt ttttttatac cattttctct tctctagctt actttattgt 15180 gagaacacaa tttataagca tataaaaggc agtatagcaa gatcccatct ctaaaaacaa 15240 atattaaaaa aaaaattagc ttggcagagg ggcaggcact tgtggtccca gctaacagag 15300 aggctgaggt gggaagattc cttgagccca gaagtctgag attgcagtga tggcaccact 15360 gcactccagc ctaggtgacc gtatgtcttc attgattatt tctgttatca gtaaggcatc 15420 tagtcaacag caggctatta gtggttaagt tttgggggag tcaaaattat acatgaattt 15480 tcaactgttc agaggtcaga gcccctaact cccagatttt tcaaggacta tctgtatata 15540 caattgtagc actgctgcta taaatgtgca gatatgtttt aaatgatcta aggctaacaa aaacatattc agcagttttt ttaattttaa gaataatcta ttcagaatgt gtatccctgt 15660 gcaattaatt acaaatagct aaaatcttgc aaaatattta atgtatcaaa tcagcatgtg 15720 atttcattta tagaaataaa agactagttt tctcccttct tttacccatg aacttggcta 15780 attaattttc tatttttccc tcaaactttt ctcaaaggtg ggtatgaagg gaaaacttgt 15840 ttttgtgaca ctagagagaa attgatgaag cctgggcaca acgtcatttg tctccctgtc 15900





acaaagttgc	aggtragctg	gccattcatc	acctggtaca	cagcacacct	catcttgacc	15960
aaagagcacc	tttcttcatc	agagaagctg	tgctattgag	agcactggga	accaggtctg	16020
aattggaaca	agaggcagaa	aagcagaatg	atrtttcttt	ctaaaatcat	ataataaaca	16080
ccaataaaat	ttctgaatag	tagtaacgtg	tcttttgtta	ttcaaagtag	tattatatag	16140
cttttcctct	tcattttatt	gaaagtttca	ggaaggggga	atgtactaaa	gcttaaaatť	16200
aatatctaat	gcttttcatc	ttgagaacag	tatgtacatc	ttgcacaact	atattttagt	16260
ttgtataaaa	tagacatata	aattacctgc	attcatggct	ttcattcatt	tatctgaata	16320
aacttactga	gcagcctcta	tgttcctggg	actgggcaag	gcattggaga	aatagataca	16380
aaagacctat	ttttggactt	ccaggggttc	ttagaccact	gtggaaaaag	ggcatacgat	16440
tacaatgcat	gttatgtaaa	aaatatggca	aagttagagt	acactggaag	catttaactc	16500
ctattgacag	aagttagttg	cacaaataga	gtggcatttt	aaatttaata	atcagatcta	16560
aattcagcca	gcaaaagaca	cagaaatttt	gtcatgtgca	aagatgggaa	atatcaaaat	16620
agaaagttgg	gggaagagtc	actttataaa	acaaatgatt	acctttagat	tgaaagaaag	16680
tcatgacccc	tattgaaaac	agacagacaa	atgtcggtat	aggatctgtt	gaggaacaga	16740
gggagtaaca	gtggggaatg	cctcaatgct	attttgcctt	tcagcagaga	tgggcagaag	16800
agatttgaaa	aagggttcat	ttttatagat	gtatgtaggt	atttgtatac	gtgtgtatgt	16860
gagttcaccc	acacatgtat	gtatgtgtat	taataaccca	ctgacctcaa	agtaagagag	16920
aaataggatt	tacccagtat	ttacccaaga	atccccatca	tccttatccc	aggattctgc	16980
agtcagcaaa	ccaggctttt	taacactctc	caatcatgag	gtatttaaag	arctccatga	17040
tttgttttgt	atttaacgct	gcaataacca	ttctcagata	aacatgtttg	agggcttctt	17100
gagatacatt	tttctaagga	aagtttatgg	gttaatagag	ctttaattgt	tttgggtata	17160
tatttccaat	tttccctacc	aacatttntg	caaaaggcac	agagcaatcc	agctgtacac	17220
atgcatgcct	gctgcatttt	catagaaagt	tcttcctacc	atgaaaacat	accttttact	17280
cagttctctg	aactgtggat	ttcacgtcag	gagagtaaaa	atgtggnngt	agttatgttc	17340
ttggttttct	ttgggagtgg	gttcttgagg	aaagagatga	caaacaagaa	taggcaactt	17400
caggtctcca	aaccncagaa	ttggcaaata	tagctaaacc	ggaaggggac	agtcattttc	17460
aagtgctgtg	tttttttaaa	atatcaggta	atttttaaaa	tctacgttta	atataacttt	17520
atcagcaatc	aaaaaaaaa	ctggtcaact	tgaatgatct	tagtcgtctt	ttctatgagg	17580
atctcttctt	gatgtaaaat	atccaaaggc	cctgttcaca	ccttaggaca	ccagttagca	17640
ttcgtttctg	ttgatcaaga	aatatatata	cagatgctct	ttgacttaca	atgaggcgcc	17700
atcctgataa	acccataata	agttgaaaat	attccaagta	aaacgtgcat	ttggtacaag	17760
tacacctaac	ctactgaaca	tcatagttag	tcctgcctac	ctgaaacgtg	cttggaacac	17820
ttacattagc	ctacagttga	gcaaaatcat	ctcacacaaa	accaatttt	ttttttgaga	17880
tgtagtttca	ctcttgttgc	ccaggctaga	gtgcagnggc	acaatctcag	ctccctgcaa	17940
cctntgcctc	ctgagttcaa	gngattctcc	tgcctnaacc	tcccaaatag	ctgggattac	18000
aggcatgtgc	caccacaccc	agctaattct	gtatttttag	tagagatggg	gtttcaccgt	18060
gtttgtcaga	caaaacctat	tttatagtga	agtgtcgaat	atctcattta	atttattgat	18120
gttctactga	gtgtatatca	cttttaaacc	atcgaaaagt	ttaaaatcat	taagtcaaac	18180
ataatctgtc	aaaccattat	agtcgaggac	catctctaat	ctgnaatatg	aaacattaca	18240
ttacatattt	atgtatcata	taatacttca	attatataaa	atgaatttac	atatgaatcg	18300
aaatagtatc	cacacatatt	taacaatata	tgtacatata	tttcatttca	catagtgtag	18360
acccctagaa	attaggtcat	tgtctgccta	ctccaaatat	catctaaata	ccatccaaag	18420
gttgaaaatt	agttecteaa	ttatgttaac	atttcagtgg	aaacattaaa	atggtagtaa	18480
ccataagctt	gccattaatc	cagtaatgac	atcagatttt	tgaagattta	tgttcaattt	18540
cagacttcat	attttaaaag	atatgagcta	tgagaacatg	acaatgatta	aaactttaca	18600
aaataaaata	tctggggaga	aaggaataat	attgggatat	attactgaga	aggaaaagaa	18660
ggtgaaagag	agacaactta	tttttcaggc	atctgtcaga	gccaagaact	ttggataaat	18720
tatttctggc	tctcacaaca	aacttgcata	aaaggttttt	taatcctaaa	tttttggagg	18780
agatacaaat	ctctgagaga	gtatgaaaat	gagtcagagt	cccataaatc	gggaacatct	18840
gttcttggtc	tcaaacccca	tgccaggacc	atgttctttc	gataacaaca	cacttaatac	18900
agtcctgctc	tcctaatcag	gaaagacaca	agggtctatg	aaatcactta	aaattcaaga	18960
tttagcctta	attagtaaaa	agaaattgca	aatttaattg	tcagaaatgt	ggaagaaatg	19020
caataacatt	gcctaggttt	atattcctca	aacctttggc	ctcagaataa	agggttaggt	19080
tccgtattgg	ccaaggatga	tcatatcaca	tttatataaa	atcgctaaca	aaatcagtca	19140





```
atattaactg aaaaatggtg ttcaggacaa tggtcaagat gaataataga cactaactcc
tattttgaca agtttataaa ttagttaagg atacaggcac atggcttaaa gacaattggc
agcacacttt tcatcccatg ttctcattct gttgttggca aataacaaat tatctttgtg
                                                                   19320
tccagtgatt tagtgatggg aaatcattct cttaaacttg ataaagaaaa ttactgttct
                                                                   19380
ccagaaaaaa atatttcaag rgagagaata tcagtgatgt aatgagaact tataaatgta
aggaatgcct cattataaac atagtattat atccagctca gtgtcaacaa tggtaatttg
                                                                   19500
tggaggaggt tagcgattct ccatgaactc gacccaatgt cataaacact catttctcaa
                                                                   19560
gaaaacattt aatttaagga accctcagtc tgctttttag caagatatgt agtttgaata
                                                                   19620
ataatattct ggtttgaact gttttaaact taggcacact gggtagttgg cctctccatg
accaactggc tttaggcatt gtgctcactt tctgtgagat ggcagtaagg gataaaactg
                                                                   19740
catectgeea geatacagag gattetggat ateatggget ateentgget tttageetge 19800
tttctggaaa taagccaaaa ataagtatct ttataaggcc tgctaggttt tatgtgcaga 19860
aattctaaat aattttcatt atgaaatata gtaaggagag gccgggcatg gtggctcacg 19920
actgtaatcc cagcatttgg ggaggccaag gtgggtggat catgtgaggt caggagttcg 19980
agaccagcct ggccaacatg gcaaaacccc atcgctgctg aaaataaaaa aaaaaaatta 20040
gctgaacaca gtggcaggtg cctgtagttc cagctgttcg gganactgag gctggaaaat
tgcttgaaac tgggaggtgg aggttgcagt gagctganat ggcatcgttg tactccagcc
taggtgacag agtgagactc tatctcaaaa atgaaaagag agagagagaa agaaagagaa
                                                                  20220
ggaaagaaag aaagaaagaa ggaaggaagg aaggaaggaa ggaaggaagg aaggaacaaa
                                                                  20280
gaagaaagaa agtgttgaat aaacttgtat ttagattgtc taggattatc tatataaggt
caaactcaat ttccagatta atattaatct attcccactt aataaaatgt ctctatttct
atatacactt aagagaccag ggattttgct gatggtaatt taaatttata tagtgaggtt
                                                                  20460
tacattttga aattaataac aatttttat tagcattaga gacttaattg tttaaagcaa
                                                                  20520
tottacctac ctactatatc agtagctaac tgctgcctac cacagcagtc acacaacagc
atgagcacaa aatgtttaca gttaccatgt ccctgcccag agggtatcca aaaattacat
                                                                  20640
tggcactaca gctaaataaa cccatgtaaa acatatacta taaaagagtg tagacatata 20700
tcacattttt atggaaagta aaacattgcc taattttaga taacaaaagg aaaattgtta 20760
tctacacaat aaatgaatta aaattctcat ttgtcataaa acagaaagga gagaaaatca
                                                                  20820
ggtcacttgt ttactagtca aaattatcag aattcaccca aatatgtagt gtgatacttc 20880
taaatacaat aatatagaga gaaatatgag ccaattacaa tcattttat ttagagtact 20940
agttatttta atgaaattta tttattgata atggaaaact ttttatttt tattttatgt 21000
tacttcaaaa taattattaa cttacggaat agcttcaaga acagtataat agactttttc 21060
ctgttttccc aaaccatatt ataataaatt gtcaacctga agcctgatca ttcagaatat
                                                                  21120
cttaagtata catttcataa aataaaagca ttctccttta tatcgatgat acagtattaa
                                                                  21180
aaaacagaaa gttatactga tattactacc atctagtcct cagatcctat ccaactttca
ctcattatcc caaaaaataa ccttgagagc aaaagaaccc aatttataat tacatgttgc
attgagttgc cttgtctctt ctgtctcctt cagtctgaaa cagttcctca gtgtgtattt
                                                                  21360
tactttaata gccttgaaac tttggatggt tacaggccag ttacatttta gaangcctcc 21420
tgatctgnct tgnctgatat ttccatatta ttagcttgag gtnacacatc tttgctaaan 21480
tatcagaaat gtttacccta tcatgtaaaa cattattta ccctgttcca gtaatgatga 21540
ccttcacttt aattgcatga atgaagtggt agcctcatgc ctggggatac cacacacac 21600
cacacacaca cacacacaca cacacaca
                                                                  21636
<210> 4
<211> 5566
```

<212> DNA

<213> Homo sapiens

<220>

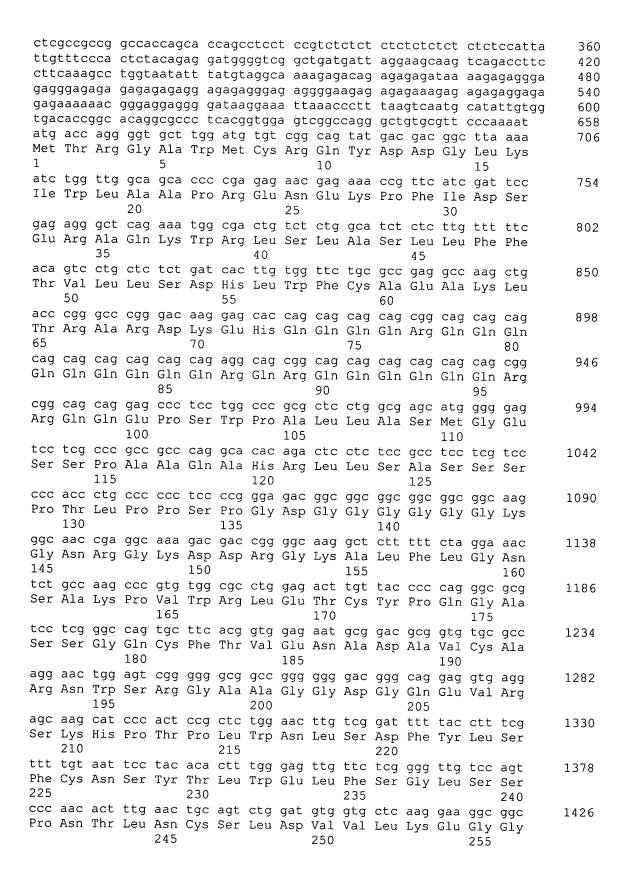
<221> CDS

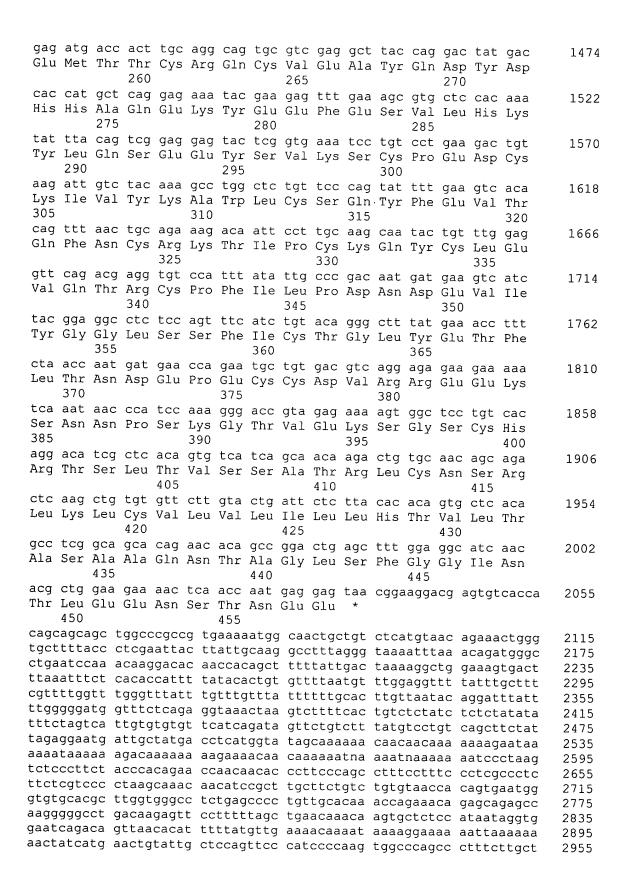
<222> 659..2032

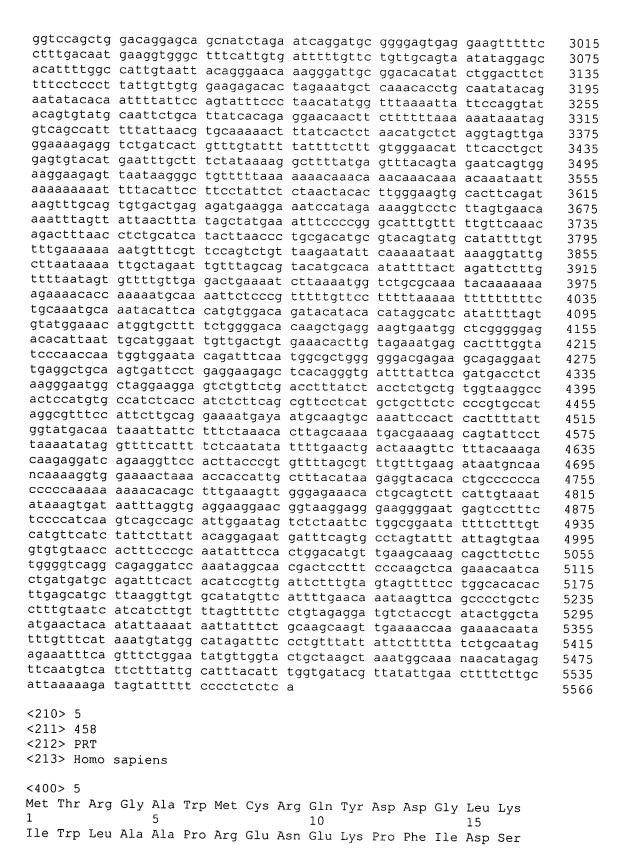
<220>

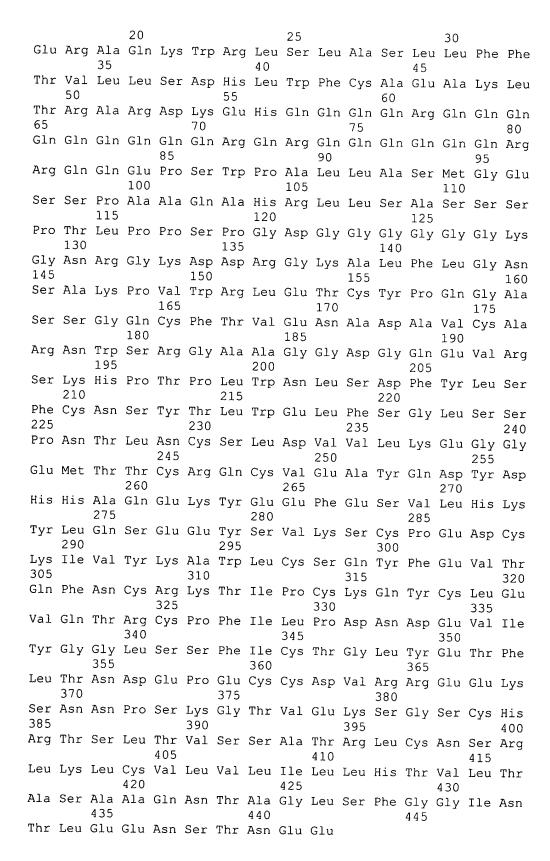
<221> allele

```
<222> 4484
<223> 99-13821-332 : polymorphic base C or T
<220>
<221> polyA_signal
<222> 2531..2536
<223> potential
<220>
<221> polyA_signal
<222> 2538..2543
<223> potential
<220>
<221> polyA_signal 
<222> 2873..2878
<223> potential
<220>
<221> polyA_signal
<222> 3307...3312
<223> potential
<220>
<221> polyA_signal
<222> 3843..3848
<223> potential
<220>
<221> polyA_signal
<222> 3859..3864
<223> potential
<220>
<221> polyA_signal
<222> 4524..4529
<223> potential
<220>
<221> polyA_signal
<222> 5536..5541
<223> ATTAAA
<220>
<221> misc_feature
<222> 10,39,2574,2580,2978,4692,4696,5466
<223> n=a, g, c or t
<400> 4
ggcagcccan gctccggact ctgggagagc cagcgcggna gcaggagcgg gaggcggagg
                                                                    60
120
ggaggaggag gggaggcgct ccgggcgccg tcagtgggca gcggaggcgc ggcatgcccc
                                                                   180
tggcagggga gagcgggctg ggctccgcgg ggccatgggg acccgcgcgc tgacaatgcc
                                                                   240
tgggtggccg gagccgcgcc agccaccctg tcacctccac catcgccccc tgcacccagc
                                                                   300
```









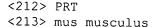
450 455

<210> 6 <211> 1791 <212> DNA <213> mus musculus

<400> 6

agccctcacg ggtggagtgg agtcggccag ggctgtgcgt tcccaaaat atg acc agg 58 Met Thr Arg ggt gct tgg atg tgt cgg cag tat gac ggc tta aaa atc tgg ttg 106 Gly Ala Trp Met Cys Arg Gln Tyr Asp Asp Gly Leu Lys Ile Trp Leu 10 gca gca ccc cgg gag aac gag aaa ccg ttc atc gat tca gag cgg gct 154 Ala Ala Pro Arg Glu Asn Glu Lys Pro Phe Ile Asp Ser Glu Arg Ala 30 cag aaa tgg cga ctg tct ctg gct tct ctc ttg ttt ttc aca gtc ctg 202 Gln Lys Trp Arg Leu Ser Leu Ala Ser Leu Leu Phe Phe Thr Val Leu 40 ctc tct gat cac ttg tgg ttc tgc gcc gag gcc aag ctg acc cgg acc 250 Leu Ser Asp His Leu Trp Phe Cys Ala Glu Ala Lys Leu Thr Arg Thr 60 298 Arg Asp Lys Glu His His Gln Gln Gln Gln Gln Gln Gln Gln Gln 75 346 90 394 Arg Gln Arg Gln Gln Gln Gln Gln Gln Glu Pro Ser Trp Pro 105 110 gcg ctc ctg gcc agc atg ggg gag tcc tcg ccc gcc gcc cag gca cac 442 Ala Leu Leu Ala Ser Met Gly Glu Ser Ser Pro Ala Ala Gln Ala His 120 125 aga ctc ctc tcc gcc tcc tcg tcc ccc acc ctg ccc ccc tcc ccg gga 490 Arg Leu Leu Ser Ala Ser Ser Ser Pro Thr Leu Pro Pro Ser Pro Gly 135 140 538 Gly Gly Gly Ser Lys Gly Asn Arg Gly Lys Asn Asn Arg Ser Arg 155 gct ctt ttt cta gga aac tct gcc aag ccg gtg tgg cgc cta gag act 586 Ala Leu Phe Leu Gly Asn Ser Ala Lys Pro Val Trp Arg Leu Glu Thr 165 170 175 tgt tac ccc cag ggc gcc tcc tcc ggc cag tgc ttc acc gtg gag agc 634 Cys Tyr Pro Gln Gly Ala Ser Ser Gly Gln Cys Phe Thr Val Glu Ser 180 185 190 gcg gac gct gtg tgc gcc agg aac tgg agt cgg ggg gcg gcc gcg ggg 682 Ala Asp Ala Val Cys Ala Arg Asn Trp Ser Arg Gly Ala Ala Gly 200 205 gag gag cag tcg tcc agg ggc tct cgg cca act ccg ctg tgg aac ttg 730 Glu Glu Gln Ser Ser Arg Gly Ser Arg Pro Thr Pro Leu Trp Asn Leu 220 tcg gat ttt tac ctt tca ttt tgt aat tcc tac aca ctt tgg gag ttg 778

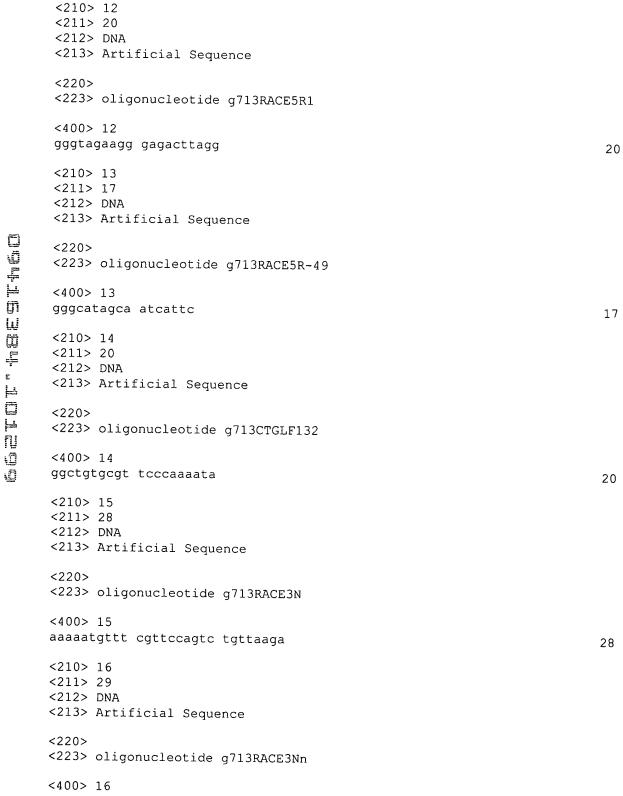
Ser As	Phe 230	Tyr	Leu	Ser	Phe	Cys 235	Asn	Ser	Tyr	Thr			Glu	Leu	
ttt tc Phe Se 24	g ggg r Gly	ctg	tcc Ser	agc Ser	ccc Pro 250	agt	act Thr	ttg Leu	aac Asn	tgc Cys 255	240 agt Ser	ctq	gac Asp	gtg Val	826
gtg cte Val Le 260	ı Thr	Glu	Gly	Gly 265	Glu	Met	Thr	Thr	Cys 270	aga Arg	Gln	Cys	Ile	Glu 275	874
gct tac Ala Ty:	r Gln	Asp	Tyr 280	Asp	His	His	Ala	Gln 285	Glu	Lys	Tyr	Glu	Glu 290	Phe	922
gaa ago Glu Sei	r Val	Leu 295	His	Lys	Tyr	Leu	Gln 300	Ser	Asp	Glu	Tyr	Ser 305	Val	Lys	970
tcc tgi Ser Cys	310	Glu	Asp	Cys	Lys	Ile 315	Val	Tyr	Lys	Ala	Trp 320	Leu	Cys	Ser	1018
cag tat Gln Ty: 325	Phe	gaa Glu	gtc Val	aca Thr	cag Gln 330	ttt Phe	aac Asn	tgc Cys	aga Arg	aag Lys 335	acc Thr	att Ile	cct Pro	tgc Cys	1066
aag caa Lys Glr 340	Tyr	Cys	Leu	Glu 345	Val	Gln	Thr	Arg	Cys 350	Pro	Phe	Ile	Leu	Pro 355	1114
gac aat Asp Asr	Asp	Glu	Val 360	Ile	Tyr	Gly	Gly	Leu 365	Ser	Ser	Phe	Ile	Cys 370	Thr	1162
ggg cto Gly Leu	tac Tyr	gaa Glu 375	acc Thr	ttc Phe	cta Leu	acc Thr	aat Asn 380	gat Asp	gaa Glu	ccc Pro	gaa Glu	tgc Cys 385	tgt Cys	gac Asp	1210
atc ago	Ser 390	Glu	Glu	Gln	Thr	Ala 395	Pro	Arg	Pro	Lys	Gly 400	Thr	Val	Asp	1258
aga aga Arg Arg 405	Asp	Ser	Cys	Pro	Arg 410	Thr	Ser	Leu	Thr	Val 415	Ser	Ser	Ala	Thr	1306
aga ctg Arg Leu 420	Cys	Pro	Gly	Arg 425	Leu	Lys	Leu	Cys	Val 430	Leu	Val	Leu	Ile	Leu 435	1354
ctc cac Leu His	Thr	Val	Leu 440	Thr	Ala	Ser	Ala	Ala 445	Gln	Asn	Ser	Thr	Gly 450	Leu	1402
ggc ctg Gly Leu	Gly	G1y 455	Leu	Pro	Thr	Leu	Glu 460	Asp	Asn	Ser	Thr	Arg 465	Glu	gac Asp	1450
tga gcg *	cagco	ag g	cgcg	tgcg	ıc ag	agcg	cagg	gct	gggc	agg	gaca	cgcg	ct		1503
tggcaca ccacccg gggcccg actcagt tcagttc	ggt g atc c ttc t	ctct ccaa caca	accc catg ccat	t tg g ac t tt	gact acag atac	tctc ccgc actg	gca agc tgt	aggc tttt ttta	ctg tgc acg	tggg cgac tttg	taac taaa gagg	at t ag g tt t	caac	aagat	1563 1623 1683 1743 1791
<210> 7 <211> 4											-				



<400> 7 Met Thr Arg Gly Ala Trp Met Cys Arg Gln Tyr Asp Asp Gly Leu Lys Ile Trp Leu Ala Ala Pro Arg Glu Asn Glu Lys Pro Phe Ile Asp Ser 25 Glu Arg Ala Gln Lys Trp Arg Leu Ser Leu Ala Ser Leu Leu Phe Phe 40 Thr Val Leu Leu Ser Asp His Leu Trp Phe Cys Ala Glu Ala Lys Leu Thr Arg Thr Arg Asp Lys Glu His His Gln Gln Gln Gln Gln Gln 70 75 90 Gln Gln Gln Arg Gln Arg Gln Gln Gln Arg Gln Gln Glu Pro 100 105 Ser Trp Pro Ala Leu Leu Ala Ser Met Gly Glu Ser Ser Pro Ala Ala 115 120 Gln Ala His Arg Leu Leu Ser Ala Ser Ser Ser Pro Thr Leu Pro Pro 135 140 Ser Pro Gly Gly Gly Gly Ser Lys Gly Asn Arg Gly Lys Asn Asn 150 Arg Ser Arg Ala Leu Phe Leu Gly Asn Ser Ala Lys Pro Val Trp Arg 170 Leu Glu Thr Cys Tyr Pro Gln Gly Ala Ser Ser Gly Gln Cys Phe Thr 180 185 Val Glu Ser Ala Asp Ala Val Cys Ala Arg Asn Trp Ser Arg Gly Ala 200 205 Ala Ala Gly Glu Glu Gln Ser Ser Arg Gly Ser Arg Pro Thr Pro Leu 215 220 Trp Asn Leu Ser Asp Phe Tyr Leu Ser Phe Cys Asn Ser Tyr Thr Leu 230 235 Trp Glu Leu Phe Ser Gly Leu Ser Ser Pro Ser Thr Leu Asn Cys Ser 245 250 Leu Asp Val Val Leu Thr Glu Gly Gly Glu Met Thr Thr Cys Arg Gln 265 Cys Ile Glu Ala Tyr Gln Asp Tyr Asp His His Ala Gln Glu Lys Tyr 280 Glu Glu Phe Glu Ser Val Leu His Lys Tyr Leu Gln Ser Asp Glu Tyr 295 300 Ser Val Lys Ser Cys Pro Glu Asp Cys Lys Ile Val Tyr Lys Ala Trp 310 315 Leu Cys Ser Gln Tyr Phe Glu Val Thr Gln Phe Asn Cys Arg Lys Thr 330 Ile Pro Cys Lys Gln Tyr Cys Leu Glu Val Gln Thr Arg Cys Pro Phe 340 345 Ile Leu Pro Asp Asn Asp Glu Val Ile Tyr Gly Gly Leu Ser Ser Phe 360 Ile Cys Thr Gly Leu Tyr Glu Thr Phe Leu Thr Asn Asp Glu Pro Glu 375 380 Cys Cys Asp Ile Arg Ser Glu Glu Gln Thr Ala Pro Arg Pro Lys Gly 395



Thr	Val	Asp	Arg	Arg 405	Asp	Ser	Cys	Pro	Arg 410	Thr	Ser	Leu	Thr	Val 415	Ser	
Ser	Ala	Thr	Arg 420	Leu	Cys	Pro	Gly	Arg 425	Leu	Lys	Leu	Cys	Val 430	Leu	Val	
Leu	Ile	Leu 435	Leu	His	Thr	Val	Leu 440	Thr	Ala	Ser	Ala	Ala 445	Gln	Asn	Ser	
	450		Gly	Leu	Gly	Gly 455		Pro	Thr	Leu	Glu 460		Asn	Ser	Thr	
Arg 465	Glu	Asp														
<210 <211)> 8 .> 23	3														
	?> DN 3> Ar		icial	l Sec	quenc	e										
<220 <223		.igor	nucle	eotic	de g7	'13LF	' 1									
<400					_											
cgct	tgct	tc t	gtct	gtgt	a ac	c										23
<210																
<211 <212																
<213	> Ar	tifi	.cial	. Sec	quenc	е										
<220					1	1275										
		rgon	iucie	000	le g7	1358	.1									
<400 gtat		gc a	gacc	attt	t aa	gatt										26
<210																
<211 <212																
<213	> Ar	tifi	cial	Seq	uenc	е										
<220 <223		igon	ucle	otid	e 71	3.LF	1.5.	1								
<400																
actg	tctg	at t	ccac	ctat	t at	ggag										26
<210 <211																
<212		A														
<213	> Ar	tifi	cial	Seq	uenc	е										
<220 <223		iaon	ucle	ot i d	o ~7	12 т	C1 F	1								
		-9011	4016	o c i u	eg/	1 J . 11.	т.э	• TII								
<4000 tgat		cc t	atta [.]	tgga	g ag	cac										25



	attgctagaa ttgtttagca gtacatgca	29
	<210> 17	
	<211> 20	
	<212> DNA	
	<213> Artificial Sequence	
	<220>	
	<223> oligonucleotide SG1polyA	
	<400> 17	
	ttttttttt tttgacagag	20
	<210> 18	
	<211> 25	
	<212> DNA	
a	<213> Artificial Sequence	
Ĩ	<220>	
Ę Li	<223> oligonucleotide SG1LR100	
(2) (2) (2) (2) (4) (4) (4) (4) (5) (4) (5) (6)	<400> 18	
<u>J</u>	tttgccattt agcttagcag tacca	25
IJ P	<210> 19	
	<211> 22	
	<212> DNA	
⇒ ¬	<213> Artificial Sequence	
÷	<220>	
U	<223> oligonucleotide g713.PU	
I A	<400> 19	
₽	aatattetta acagaetgga ac	22
	<210> 20	
	<210	
	<2112 ZZ <212> DNA	
	<213> Artificial Sequence	
	<220>	
	<223> oligonucleotide g713.RP	
	<400> 20	
	ctttatagct atgaaatttc cc	22
	<210> 21	
	<211> 24	
	<212> DNA	
	<213> Artificial Sequence	
	<220>	
	<223> oligonucleotide g34301.PU	

<400> 21 ctgatcactt gtggttctgc gccg	24
<210> 22 <211> 22 <212> DNA <213> Artificial Sequence	
<220> <223> oligonucleotide g34301.RP	
<400> 22 aggactecce catgetegee ag	22
<210> 23 <211> 23 <212> DNA <213> Artificial Sequence	
<220> <223> oligonucleotide SG1LR1102	
<400> 23 . aaaatactgg gaacagagcc agg	23
<210> 24 <211> 18 <212> DNA <213> Artificial Sequence	
<220> <223> oligonucleotide SG1LF790	
<400> 24 gcacttagag cgcggggt	18
<210> 25 <211> 15 <212> DNA <213> Artificial Sequence	
<220> <223> oligonucleotide SG1LF834	
<400> 25 gccggaggca gccca	15
<210> 26 <211> 17 <212> DNA <213> Artificial Sequence	10
<220> <223> oligonucleotide moCTGR1511	

Ø
<u></u>
 - -
m
ليا
M
Ļ
Đ
⊨
<u> </u>
ΓU
Ū
Ū

	<400> 26 tgtcctcgag cgtgggg	
		17
	<210> 27	
	<211> 27 <212> DNA	
	<213> Artificial Sequence	
	<220>	
	<223> oligonucleotide moCTGLR20	
	<400> 27	
	cggaggaggg gatacggaaa ttaaacc	27
	<210> 28	
[77]	<211> 25	
. M	<212> DNA	
# #	<213> Artificial Sequence	
	<220>	
() 	<223> oligonucleotide moCTG1440	
w M	<400> 28	
-	tgggtcactg ctgctctgtg ccaag	25
E .	<210> 29	
F255	<211> 20	
	<212> DNA	
 71	<213> Artificial Sequence	
	<220>	
Ī	<223> oligonucleotide moCTG5RACE1	
	<400> 29	
	tcacagtgtc ctcggccact	20
		20
	<210> 30	
	<211> 20	
	<212> DNA	
	<213> Artificial Sequence	
	<220>	
	<223> oligonucleotide moCTG5RACEn	
	<400> 30	
	tcctccacac agtgctcacg	20
	<210> 31	
	<211> 983	
	<212> DNA	
	<213> Homo sapiens	
	<220>	





120

180

240

300

360

420

480

540

600

660

720

780

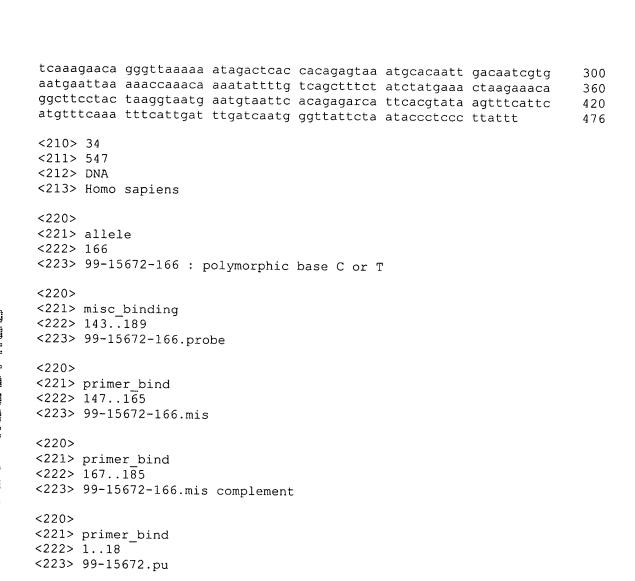
840

900

960

```
<221> misc feature
<222> 14
<223> n=a, g, c or t
<400> 31
aaaaaaaaa aaanaacaaa aacaaaacat gaaacaggaa gacaggaaag atccatttga
cagagtactt gacacaggag agagaagaaa tactcatgta tctgaaagta ttcaaagggg
gagtgttagg agatgaatta atttaaaaaa tgagtaagag taaaatagtt taaagttaga
ccctgaggaa ctccagggaa gacaaagtaa cacaaggaac aagcaatgtt agccactgcc
taactttcct cagggtcatg tgtgcctcgc cataattatg taaacactta cattgttaaa
acgaaattcg gagaactagt ttgagtaaag gggaaaagaa agtatgttat tcatgtcgga
gttggaaata tgtgataggt tgaaattctc aatttcctaa ttggaaatca ttaagtcata
ctgaaacctg aaaattcaag aactgacaac acaattgatg ttgagatatg gaatttggta
cctgatgaaa gattagaaaa ttattaaaag caatttcttc tgggtggtgc tacaagatgg
aagaagaaag gacagaaagc tetteataat caggtagaeg etttgaettt ttaagtggta
tgcctatatg cctttaaaaa acaactcaat ttaaaagaaa attaagagat gctaacagcc
gatttaaaga aaatttagta aaatattcaa ttgtataaag atacacaaaa tattggttat
ctacatgata gcaaagatga attaagggat ggggataaaa ctcttctcaa taacaccaaa
attaaaataa aacataattc atatattag aaatatcatt acagaaatat gttgaacttg
tattaacagc ctctcctcaa aggtagcatg gagaatcatg caaacttaat ttggagatac
aaaaaaaatt gagaatgtgt agtgttgttc tttaattcta actgtaatgg ctgaataata
ttttgatcat gattgtgata cta
<210> 32
<211> 450
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 298
<223> 99-15663-298 : polymorphic base C or T
<220>
<221> misc_binding
<222> 275..321
<223> 99-15663-298.probe
<220>
<221> primer bind
<222> 279..297
<223> 99-15663-298.mis
<220>
<221> primer bind
<222> 299..317
<223> 99-15663-298.mis complement
<220>
<221> primer bind
<222> 1..18
<223> 99-15663.pu
<220>
```

```
<221> primer bind
 <222> 430..450
<223> 99-15663.rp complement
<400> 32
tcccaccttc ttctaaacgt gttgcttcaa tacgttgata ggtgaggaca cttaaaaatt
                                                                         60
agactttata gaaataggtt tttttttgtt tacatatata gttcttttgg tatcatatat
                                                                        120
ttagcctctt tctaaaattt atttttgat actgaaggga gaaataggga gttattaatc
                                                                        180
aacaggcatt aattttagtc aagcaaaata aataagctgt agcgatctgc tctgtaacat
                                                                        240
tgtacctaca gccaacaatt atatgttgtc cacttaaaaa tgtgttagat ctcatagyaa
                                                                        300
ctcttcttac cacaataaag taaaaattct gaaacaataa gtgaatacct aaataataca
                                                                        360
aacaaataca atattgtagt tttgggcact taataaatga cagcctcatt tctcaattag
                                                                        420
agatcatcac aagttagaca gatgacgatg
                                                                        450
<210> 33
<211> 476
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 398
<223> 99-15665-398 : polymorphic base A or G
<220>
<221> misc_binding
<222> 375..421
<223> 99-15665-398.probe
<220>
<221> primer bind
<222> 379..397
<223> 99-15665-398.mis
<220>
<221> primer bind
\langle 222 \rangle 399..4\overline{17}
<223> 99-15665-398.mis complement
<220>
<221> primer_bind
<222> 1..20
<223> 99-15665.pu
<220>
<221> primer bind
<222> 458..476
<223> 99-15665.rp complement
<400> 33
cgtaaatgtg aaaagcatag cctcttcttg gaatgttaag tataaatatc tgaaatactg
                                                                        60
ggcttgatat gtcaacagga gattgatgga taaaaataga attttatata aaaaacaact
                                                                       120
ggacatatta gattgttaac ttggaagaaa gaccatattc aaagaagaaa acatagtgac
                                                                       180
taatttcaaa catttaaagt cttccctgtg gaaacaaagg aatatctttg ttctaacact
```



tgattattat tatcagtgtt attattatcc taatcctaag taatccaata aaagaaaaat 120 acatctgtgc ctgtgcgtat gtgcacgtgt gtgcagtcaa atacaygttg agtaaaggta 180 aagtctagct gtatttaatc aacctacctg aatcctcagg aaaaaattct aaacctagtt 240 taaaacatgt aaactctaag ctctctcctt atagtcagtt agtagcagca catcttaaaa 300 tctggtgtga atattctctt agttctacat gagtctaact aaacagagga ttattcttag 360 gtgtttgaaa gagacatatg tgacactgct gttttgagaa caatttaagt gttgtcttgt 420 catgtacaga agttctcata ttactttaca taaatggttg cataattgtt ttatagtaaa 480 taatagactg tcaatatttc taggataact ccaaaacaaa atttcctaga mmacattttg 540 aaaaggg 547

60

ccaataccat aactcctcta taggacatgg aagagtatta tatatgacaa atgattgcta

<210> 35 <211> 502

<220>

<400> 34

<221> primer_bind <222> 533..551

<223> 99-15672.rp complement

```
<212> DNA
 <213> Homo sapiens
 <220>
 <221> allele
 <222> 185
<223> 99-15664-185 : polymorphic base G or T
<220>
<221> misc binding
<222> 162..208
<223> 99-15664-185.probe
<220>
<221> primer bind
<222> 166..184
<223> 99-15664-185.mis
<220>
<221> primer bind
<222> 186..204
<223> 99-15664-185.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-15664.pu
<220>
<221> primer bind
<222> 483..502
<223> 99-15664.rp complement
<220>
<221> misc_feature
<222> 54
<223> n=a, g, c or t
<400> 35
gtttaccatt agcactgtca tatttgtgtg acttgtcatt ctctacagcg gagnacgggc
                                                                        60
tggcacgggg cctgatgctg acttgcacaa gggaagcctc ctgtctctga cttccccagg
                                                                       120
ataattcctg gggaaagtgt gctccctagt gttaagagcg gtttaatggc tggagggttt
                                                                       180
cagckggctg accaggcaga gaaggaggt gaatcacctc tcagcactct ccacttagac
                                                                       240
tttgtgtggt cgtcgggtgg tcaaaccttc taactagttg tattgcagat ttggcattcc
                                                                       300
agtgcaaaca aaagacagaa acacaatgtt cacatgcttt ccagagatca cctggatatc
                                                                       360
agatcatttg attttcaagt aagtcgaaac cttggtggaa atcattaact atcctgttta
                                                                       420
tgaccaaaaa ataaaatccc aaatttcttc tcttcatttc ttacctgctt taaaattgta
                                                                       480
tccaaagcgt graagtaaaa ga
                                                                       502
<210> 36
<211> 455
<212> DNA
<213> Homo sapiens
```

```
<220>
      <221> allele
      <222> 205
      <223> 99-5919-215 : polymorphic base A or G
      <220>
      <221> misc binding
      <222> 182..228
      <223> 99-5919-215.probe
      <220>
      <221> primer bind
      <222> 186..204
      <223> 99-5919-215.mis
      <220>
      <221> primer_bind
<222> 206..224
      <223> 99-5919-215.mis complement
     <220>
     <221> primer bind
     <222> 1..19
     <223> 99-5919.pu
     <220>
     <221> primer bind
     <222> 435..455
     <223> 99-5919.rp complement
     <400> 36
     ctacagcaat gcagatttca attctgccat tgaattccca gacatattcg tcatccccat
                                                                             60
     tttcatcccc caccacctg ccattttctt cgtgttaact tgttttcctg actcacagaa
                                                                            120
     atcacctttt cctgtataca tttttaggat gtcagacttt attctaatga tttctcctag
                                                                            180
     ttgcccccca aaattgtatt ctacrgtgtg attttaaagc tgaattttca agatgatatt
                                                                            240
     tcatatctat attttcacaa gcttttcttc tatgaatgtt attgtcagct gtcagggtgt
                                                                            300
     gagatggtac ttgatactac attctttcca agctgttgcc tgaatcggtt taagacaaag
                                                                            360
     tcattactag gctgtaaact gttgctctgc aaaattgagc agcacgtatt taaccactca
                                                                            420
     tacttcttag ctctccaaca ctttgagtca ataga
                                                                            455
     <210> 37
     <211> 450
     <212> DNA
     <213> Homo sapiens
     <220>
     <221> allele
     <222> 157
     <223> 99-5862-167 : polymorphic base C or T
     <220>
     <221> misc_binding
     <222> 134..180
     <223> 99-5862-167.probe
```





120

180

240

300

360

420

```
<220>
<221> primer bind
\langle 222 \rangle 138...1\overline{5}6
<223> 99-5862-167.mis
<220>
<221> primer bind
<222> 158..176
<223> 99-5862-167.mis complement
<220>
<221> primer bind
<222> 1..20
<223> 99-5862.pu
<220>
<221> primer_bind
<222> 430..450
<223> 99-5862.rp complement
<400> 37
aatcaaggta gagatgtatg agaaatagcc ggttaaagaa acagcattac tttcagacta
tcttttattt gaaatacacg tggggaaacc agaaggtgaa accccttagg agatggatat
aggatactaa aatctgagtt agaaaaattt gagcatyagc accttacgtg tcatgctaag
atagtgaatg agactgcaca ggaattgcat gcagtttaac ggaaaaagaa gtcgaaagat
aaattcctag aacactaaca ccgagttatg ggaggagaaa tatcctgcac aggtcactct
gggagacatg tcaattgttt agccaatatc catttaactc atctttcttc ctaatgaaaa
ccgaatttgg agaagcaggt agtgcccctg gctagaaata tgaaccttcc cagcttctct
catgcactga actgacaaag ttcaggtctg
<210> 38
<211> 403
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 292
<223> 99-16032-292 : polymorphic base A or C
<220>
<221> misc_binding
<222> 269..315
<223> 99-16032-292.probe
<220>
<221> primer bind
<222> 273..291
<223> 99-16032-292.mis
<220>
<221> primer bind
<222> 293..311
```



```
<223> 99-16032-292.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-16032.pu
<220>
<221> primer bind
<222> 384..403
<223> 99-16032.rp complement
<400> 38
gttgttaccc cacttettcc ccccagetcc cccttectca cacagttcat gccacatgcc
                                                                        60
actotcctgg actactggaa atgcgtcagt ccactctggg ctcatcccat catccccat
                                                                       120
gctgcaacct gagagagat tgcaagttgc aaatctgatc ttgtcaccac cactctccac
                                                                       180
actaaatccc tctaatgcct ccccctttct ttttggataa attccttctg cttgcatagc
                                                                       240
cacgtggttg gcttctatag catcacttca cactgtggtc acctgccttc tmctcactca
                                                                      300
ggaacttctc tccattgaag aagttcttct tccccatctc cagggctttc ccactgacag
                                                                      360
403
<210> 39
<211> 476
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 118
<223> 99-16038-118 : polymorphic base A or G
<220>
<221> misc binding
\langle 222 \rangle 95..\overline{1}41
<223> 99-16038-118.probe
<220>
<221> primer_bind
<222> 99..117
<223> 99-16038-118.mis
<220>
<221> primer bind
<222> 119..137
<223> 99-16038-118.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-16038.pu
<220>
<221> primer bind
\langle 222 \rangle 456...476
```

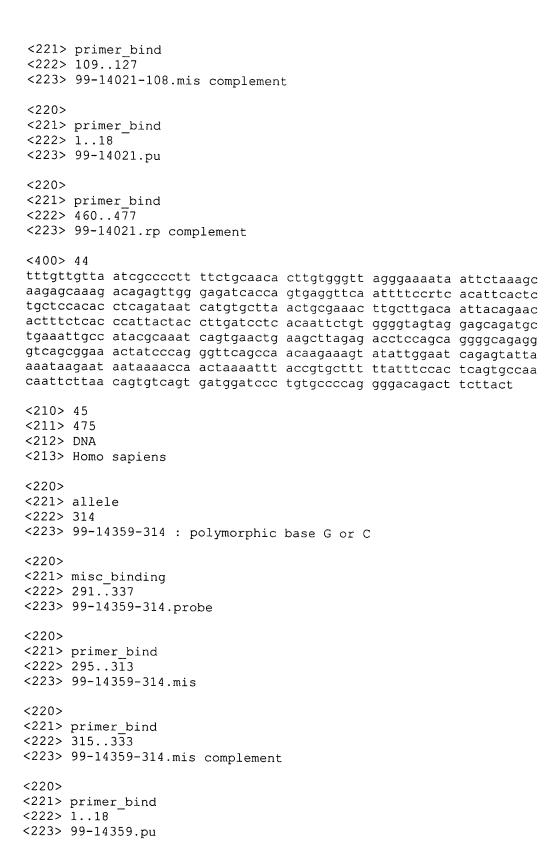
```
<223> 99-16038.rp complement
<400> 39
gttgcttatt ctttctctct tctgcagggt ataaaggaat ctgaacacga ctgatatttt
                                                                        60
ctttaatttt tagatccaga tatacattgg gtaaaatcta cttcataggt tttcaaarga
                                                                       120
gcattcttct gagcaaatct gaaaactctc taaactctat tggtatgtta ctctttatct
                                                                       180
ttatatgaat ttaaattctt ctagaagtta gataaaactg tggtaaagct acataatact
                                                                       240
tttgacatat tttcaagcgt agacaaactt caattaattt gtaagataca ggaagaaaat
                                                                       300
ttttccagtt aaaatgtacc tcttggtttc tggagtgtta gcaaccattc acacttacag
                                                                       360
ttcaaacagt gcaaccttgt aaaacatata taacttatga agagatcgat atctctttt
                                                                       420
ataaagcaaa caagtaaatt tttccctcaa tccatgattt atttttgtga agtggg
                                                                       476
<210> 40
<211> 498
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 133
<223> 99-5897-143 : polymorphic base A or C
<220>
<221> misc binding
<222> 110..156
<223> 99-5897-143.probe
<220>
<221> primer bind
<222> 114..132
<223> 99-5897-143.mis
<220>
<221> primer bind
<222> 134..152
<223> 99-5897-143.mis complement
<220>
<221> primer bind
<222> 1..18
<223> 99-5897.pu
<220>
<221> primer bind
<222> 475..492
<223> 99-5897.rp complement
<400> 40
aaaagtgttt gccagtcctg tttcttacag agcacagaac tcagatgctc ttataaagat
                                                                       60
acaggataaa tcacatcatt tcctgctcca tcatcagaat attattatat gatttagatc
                                                                      120
acttttttaa aamagaacat ggacttagta cagaacaaca gcaaaagcct ggggaaggag
                                                                      180
aggagtgcac catgaggagt caatggggag cagaagccag tccatttgac tgatttggtt
                                                                      240
cgtgtgcaaa ataattgcta aataattgca tatatgtgag actccgggta ttttcaaaac
                                                                      300
cagctggcaa aattgtgtta ttctctaccc tctgctggct ttcacgggtt ctctgttctc
                                                                      360
```

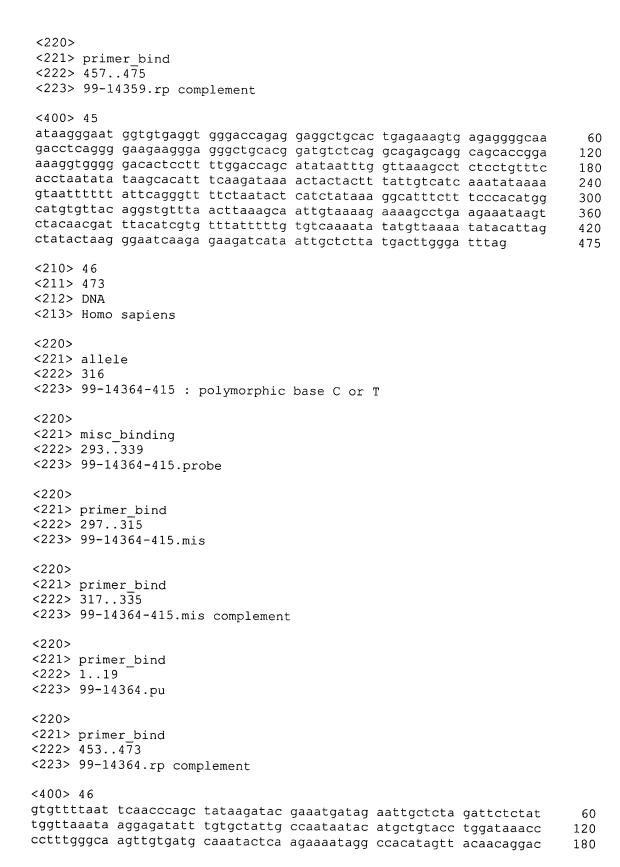
<213> Homo sapiens

```
tctccttttc ctccattctc ctcttaccct aattcctgac cactgtaatc caataatcta
                                                                       420
aggttttagg atttggatga ctaaggttac ccatggaatt gtttggaaat gtagacctgt
                                                                       480
aatggagagg ggagaaaa
                                                                       498
<210> 41
<211> 517
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 360
<223> 99-13601-360 : polymorphic base A or G
<220>
<221> misc binding
<222> 337..383
<223> 99-13601-360.probe
<220>
<221> primer_bind
<222> 341..359
<223> 99-13601-360.mis
<220>
<221> primer bind
<222> 361..379
<223> 99-13601-360.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-13601.pu
<220>
<221> primer bind
<222> 500..517
<223> 99-13601.rp complement
<400> 41
gttttacttg acagttacca agaattgttt cgcatttaag aaaattatat ctttgatggt
                                                                       60
tccctcatta atggtgcctg gatacccaat gcaacacac tacatcaaac tgcatttgta
                                                                      120
actgttggat tcataatgat tctacctaag atgcaagcat acggcatcat tgtgccttgt
                                                                      180
tgtatggata tgcttgagaa gtcacatgct gaaatacata tattttaaat ttgacagtat
                                                                      240
ctcctacaat attttcttta tattatagta aggtattaca ttacagttta aaacttatga
                                                                      300
ctataagcag gtgatattat ctatgaattt catgtgaaat tagcaaaggg acagtctcar
                                                                      360
atgtttgctg tataaagtgt atttgaagcc tgatagggtt gagaaacact cagctacagt
                                                                      420
aagtaaaaac agctctctta gtggttgcct tgttgagaag atcttgaaaa caaggttgaa
                                                                      480
aatacaaaag aaactgtgtg gagtctacaa agatatt
                                                                      517
<210> 42
<211> 533
<212> DNA
```

```
<220>
<221> allele
<222> 97
<223> 99-13925-97 : polymorphic base A or G
<220>
<221> misc_binding
<222> 74..120
<223> 99-13925-97.probe
<220>
<221> primer_bind
<222> 78..96
<223> 99-13925-97.mis
<220>
<221> primer bind
<222> 98..116
<223> 99-13925-97.mis complement
<220>
<221> primer_bind
<222> 1..20
<223> 99-13925.pu
<220>
<221> primer bind
<222> 513..533
<223> 99-13925.rp complement
<400> 42
catggaagta aaagcatatc ttcattataa gacttctaca caaattatca catctttact
                                                                        60
tacagcagct gaaacctgga aacaactcta atgcccrtca acagaggaat ggatggataa
                                                                       120
agaaactgtg atgcagtgga atacgactca acgaagatga gactaaaaat aattatactg
                                                                       180
agtaaaagaa tccaaacaaa atagagcaaa cactgtgcca tcctgtttat accttactcc
                                                                       240
agtaaatgca aactaataca caatgaaaaa aattacttat ttgagaactg gggagaggaa
                                                                       300
ggagagggaa aggggtagat aaagaaaaga ggagagatta aaaggagcat aagaaaacct
                                                                       360
cagagaataa taggtttgtg gtaaacatta ccgtggtaat gtttttaggg tatattcaca
                                                                       420
tgtaaaaact tatccaatta tacattttaa atatgtacag tttagtgtgt cagttatgcc
                                                                       480
tctgtaaagt tgattttaaa aaaagtccta ttccaagtym acaatttcat ttg
                                                                       533
<210> 43
<211> 480
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 201
<223> 99-13929-201 : polymorphic base A or C
<220>
<221> misc binding
```

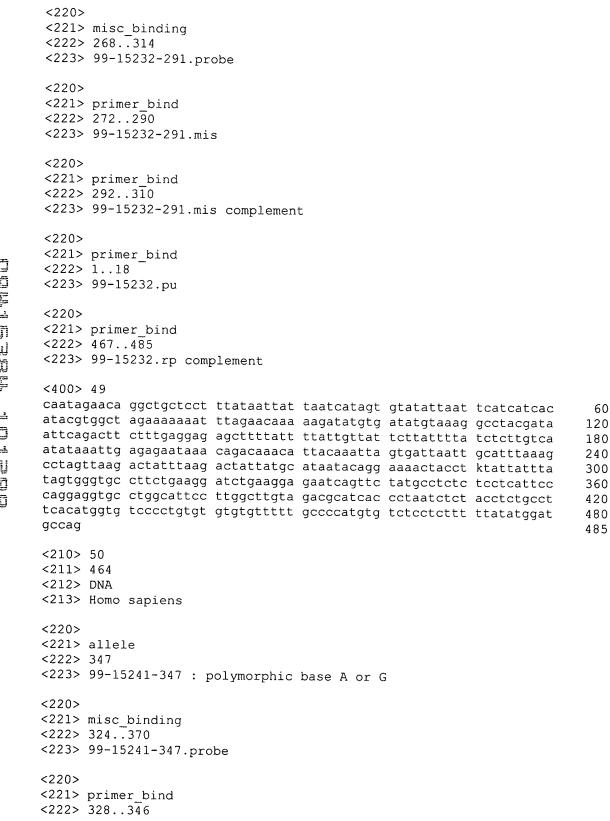
```
<222> 178..224
<223> 99-13929-201.probe
<220>
<221> primer bind
<222> 182..200
<223> 99-13929-201.mis
<220>
<221> primer_bind
<222> 202..220
<223> 99-13929-201.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-13929.pu
<220>
<221> primer bind
<222> 460..480
<223> 99-13929.rp complement
gggagaatac taataatgga agcattactt ttattttttc tataaattcc tctggaaata
                                                                        60
tgtatttctt atgtcctaag gttattaaca aaaagagaaa ataatttctg atttataatt
                                                                       120
cactttcctt caaaaaataa taactcagtg tctagtaagg taaagcaaaa aaagttaaaa
                                                                       180
gaacccataa gtttatttta maatacctac tcagaagcaa aactgacttt ctattaaaaa
                                                                       240
ttaaaaaaaa aagttttctt attattgttt tgtttccttg tttttaggtg atgggattgt
                                                                       300
atttgcaact ctctggtcag taagtgataa aatgccattt ctatgcaccc acctggcctg
                                                                       360
tgtgactggg agaatctctc tttttattaa atgtgcttca agttttaaca actgactttt
                                                                       420
gttagtgata tgatttatct acccgtgact gtcaaacaac acagatgatt tgcatatctc
                                                                       480
<210> 44
<211> 477
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 108
<223> 99-14021-108 : polymorphic base A or G
<220>
<221> misc binding
<222> 85..131
<223> 99-14021-108.probe
<220>
<221> primer bind
<222> 89..107
<223> 99-14021-108.mis
<220>
```





```
ttacctaatt ccccatggtc atttggctga ttcagtcagt tgctttcaag cctaggttct
                                                                        240
 tggctcaata ttattacata aactagaatt ttcctattac tattaatttt actttgtatt
                                                                        300
 tttctttata aacttygtac ttattgcttg tcaaatttca gcagaagtac aactcctgag
                                                                       360
agaataatgc tggctcagag ttttgagatg ataacccttg tctatgaaac tgatgaagtt
                                                                       420
ggacttaaca acgaacactc cccacagaac tcctgatgct caaatgtggc taa
                                                                       473
<210> 47
<211> 502
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 99
<223> 99-15056-99 : polymorphic base C or T
<220>
<221> misc_binding
<222> 76..122
<223> 99-15056-99.probe
<220>
<221> primer bind
<222> 80..98
<223> 99-15056-99.mis
<220>
<221> primer_bind
<222> 100..118
<223> 99-15056-99.mis complement
<220>
<221> primer_bind
<222> 1..18
<223> 99-15056.pu
<220>
<221> primer bind
<222> 482..502
<223> 99-15056.rp complement
<400> 47
caggaaactc acaagaagsc agatttcctt cgagcacctc ctgaataaag aggcaaaggc
                                                                        60
cttcttaact cttacaattt acaagtggct atgagtgcyt ttatagttcc cataataatt
                                                                      120
tctccacgta gacttcctaa ataataattt ctcctgtttt atattctctg tgcttatgtt
                                                                      180
tatatcaaac aagttaccac ttaatcaaat gccgatttgc attgctcact atgtaacttt
                                                                      240
aattttcttt gcctcttatt tttggatctt aattctaaaa ctagatgatc ataaattcat
                                                                      300
ttaggaataa gcttgtgatc tagccttctt ttgaacccct ttgtgctcct cacaatattt
                                                                      360
gtttcgatga aacagtgagc aacatttgat ctatgattgt taatagaaaa acaccaatgt
                                                                      420
ctcaagttat tgtaaacata ggcataattg acctttggtt ctataaatat gtttggtgtt
                                                                      480
ccccaaaata cgtctccctt tt
                                                                      502
<210> 48
<211> 494
```

```
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 412
<223> 99-15229-412 : polymorphic base A or G
<220>
<221> misc binding
<222> 389..435
<223> 99-15229-412.probe
<220>
<221> primer bind
<222> 393..411
<223> 99-15229-412.mis
<220>
<221> primer bind
<222> 413..431
<223> 99-15229-412.mis complement
<220>
<221> primer_bind
<222> 1..20
<223> 99-15229.pu
<220>
<221> primer_bind
<222> 476..494
<223> 99-15229.rp complement
<400> 48
ctgtcattga gaaatgctac caataatact tagagaattt gatacaactc agtctgaaaa
                                                                        60
agctaagatt agcagaacag agctgtctcc aaatatttga agaactattt tatttaaggg
                                                                       120
attggaccca tttttgtatg tagttccaga ggagcagatg gtgaccactg tccaggcaga
                                                                       180
tgtgtctcaa tgtaaggaca acatctgtaa tattaataat tagaatgtat cctgtaattt
                                                                       240
tetetetace ettggaaace agtegagate cagagtettt caetgggagg ettaaageet
                                                                       300
agagcagcct tggtgctaga ggcggacagg gataatgaac taatcttgaa ccaattcatc
                                                                       360
catagcaatc tcaatgcttt cgttagctct tataggtatt taatacggcc avaggaatga
                                                                       420
aggtagtett getggtttag aageeetgee taccacaace cetacaccae eccateceet
                                                                       480
gcatagtctg atgt
                                                                       494
<210> 49
<211> 485
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 291
<223> 99-15232-291 : polymorphic base G or T
```





```
<223> 99-15241-347.mis
<220>
<221> primer bind
<222> 348..366
<223> 99-15241-347.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-15241.pu
<220>
<221> primer bind
<222> 444..464
<223> 99-15241.rp complement
<400> 50
gttatgggtt gaaaatctct gagttcttgt acatacaaaa attttactgt tgtcacagtt
gaatcttagt ttagatgggt ataggatttt tattcaaaat gcttttactc cataagttta
                                                                       120
aaaatattgt tacattttcc tcaagtatct gatgttattg atgagaagtt taattctaat
                                                                       180
ttgactcttg ttcccttgta ggtactattt gttttccagt ttgggaagct tacatttctt
                                                                       240
aaaattcaca acatataatt tacatactac acaattcttt ttaaagtata caattcaatg
                                                                       300
catttagtat gttttagtac atataactta aattatgtat atacaaratc tctttataat
                                                                       360
atttgtagaa tatgtagcat attcacaaga ttgttcaacc atcaccactc tctatttcca
                                                                       420
gaatcttttc ctccaaaaag aaaccctgaa cactatgatg aata
                                                                       464
<210> 51
<211> 550
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 196
<223> 99-15244-196 : polymorphic base A or G
<220>
<221> misc_binding
<222> 173..219
<223> 99-15244-196.probe
<220>
<221> primer bind
<222> 177..195
<223> 99-15244-196.mis
<220>
<221> primer_bind
<222> 197..215
<223> 99-15244-196.mis complement
<220>
<221> primer_bind
```



120

180

240

300

360

420

480

540

```
<222> 1..20
<223> 99-15244.pu
<220>
<221> primer bind
<222> 532..550
<223> 99-15244.rp complement
<400> 51
ctgcttctgg ttatgttttc ctaattgcca aaatggtaaa aatgagaata atcattgaaa
gagaaagcat aaagtagcaa aaatcctttc cagattaaaa aacgaagcaa agcatgtttc
ccaagtaata atacteteat ettecteeet aateetttae eccaetacea gaagaagagt
aaaatgtccg gatatrtttg aaggtaaaga tttctccttt taataaaatt agtcaccttg
tacacatcag tagatcttga gaatgaaaag cttttctagt acattcattt caacctataa
atgtttgact tttctctgtc attcatttac gacctgtgat cttttcattc cctttcagtt
agaatatttt tcaaattttt attgatattt tctatttaac ccataggtta tttggaaata
cattgtttaa tttctaatat atttgctttt ttttctactt atttctttt ttcttaattc
cacactggtc caaatatatt ctgcatatga tttaatattt taagttctgt agagactaac
cttgtgccct
<210> 52
<211> 452
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 404
<223> 99-15252-404 : polymorphic base C or T
<220>
<221> misc_binding
<222> 381..427
<223> 99-15252-404.probe
<220>
<221> primer bind
<222> 385..403
<223> 99-15252-404.mis
<220>
<221> primer bind
<222> 405..423
<223> 99-15252-404.mis complement
<220>
<221> primer bind
<222> 1..18
<223> 99-15252.pu
<220>
<221> primer bind
<222> 433..452
<223> 99-15252.rp complement
```

```
<400> 52
atgggggcat atagcaaccc tttagaaaca aaactacaaa aggtaagctt gtcttcttgc
                                                                         60
atttcctttc tcttactaca tttaacatgg gaggttttct atgtctcaca ttcaaatatt
                                                                        120
ctcactcggg ctgcctaatt tttccctgat tttccatcac tctttatgaa ggcttgctac
                                                                        180
tttagaatac acattttctt aacagaagat aataatcaga agatgtctcc caaatataag
                                                                        240
tccaaatctt tcctatcatg ctgtgttctt tggctctttt gactttattt gaagtcagcc
                                                                        300
ttgaagggga tagagatagg ctgtatgaag tccacgctga gaagttttgc cctgcctac
                                                                        360
ttgtcctgta atatttcatg gatagcccag tggtgattaa accygtgtgt acaggaataa
                                                                        420
ccatgagaat ttgttaaaaa tataggctct gg
                                                                        452
<210> 53
<211> 477
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 382
\langle 223 \rangle 99-15253-382 : polymorphic base C or T
<220>
<221> misc_binding
<222> 359..405
<223> 99-15253-382.probe
<220>
<221> primer bind
<222> 363..381
<223> 99-15253-382.mis
<220>
<221> primer bind
<222> 383..401
<223> 99-15253-382.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-15253.pu
<220>
<221> primer bind
\langle 222 \rangle \ 459...477
<223> 99-15253.rp complement
<400> 53
aaaatcaatt ccccaacact cattttgtac gctaattttg taagatcctg aaaagtttca
                                                                         60
ctattttatg gtttcatgtg ttacagatga aaaaaaaact agaattcaaa ttttctgagt
                                                                        120
ttttttttac aatattttat gattacaaag ttagaagact aagaataaaa tggcctaatt
                                                                        180
tccataatgt gagtggtaaa tgcagagcac tggcctaaag aaaatatttc aaaaaattag
                                                                        240
tcatcttttc cttaattttt ttccaaccta tgatctgttg aatgagcatt ttgcatatat
                                                                        300
aaataaataa attactttgt aaataatctt gactggtttc tgttgaccac agtaacccac
                                                                        360
tgcacagcac agcctgtaat tyctatgaac ctagggaaat gtatttaagt ttatttttg
                                                                        420
```

```
attacacagg teeteattgt gtaactaaac attgeataga atatgeeagt gatgatg
                                                                            477
      <210> 54
      <211> 456
      <212> DNA
      <213> Homo sapiens
      <220>
      <221> allele
      <222> 392
      <223> 99-15256-392 : polymorphic base C or T
      <220>
      <221> misc binding
      <222> 369..415
      <223> 99-15256-392.probe
      <220>
      <221> primer bind
      <222> 373..391
      <223> 99-15256-392.mis
      <220>
      <221> primer_bind
      <222> 393..411
      <223> 99-15256-392.mis complement
      <220>
      <221> primer bind
      <222> 1..18
      <223> 99-15256.pu
      <220>
Ū
      <221> primer bind
      <222> 439..456
      <223> 99-15256.rp complement
      <400> 54
     cctctctatg atgcttccta ttaagcaatt ggggaaatgt aataaacaag ggttggtgag
                                                                             60
     catcttcctt agtgagatgt ttttggaaga attggataat tgagtgaata atagtgagaa
                                                                            120
     actcctgtgt ctgatgttgc tccatgttgg aatgctttta tgttctcaga gaatgagtca
                                                                            180
     ctgagagcca attgtgatga tacacaatgg ttttacccag gttggatatg gtcctctgta
                                                                            240
     ctggtaccct ttaagtcagt ggcactaatc agtcagtcat tgtcatgctt tgtgttggtc
                                                                            300
     catcatatgg tatgccctct tagagaacat cctgattagt ccttagacat cttttcaatt
                                                                            360
     tgaacactgg ggctcctcat tcgggtaaaa aytatggaca gtcagtgaaa ctgttgcaat
                                                                            420
     ggcccctcat agcagattgg atctcaatgc actttg
                                                                            456
     <210> 55
     <211> 501
     <212> DNA
     <213> Homo sapiens
     <220>
     <221> allele
```

```
<222> 200
<223> 99-15261-202 : polymorphic base A or G
<220>
<221> misc_binding
<222> 177..223
<223> 99-15261-202.probe
<220>
<221> primer bind
<222> 181..199
<223> 99-15261-202.mis
<220>
<221> primer bind
<222> 201..219
<223> 99-15261-202.mis complement
<220>
<221> primer_bind
<222> 1..19
<223> 99-15261.pu
<220>
<221> primer bind
<222> 481..501
<223> 99-15261.rp complement
<400> 55
cttctaatcc tttgtttcca cttattttat ttcattcctc attttatccc ttttttctaa
                                                                        60
attccatttt attatactta aggtgctttt aatatggtta tcatactcct gatagtgtta
                                                                       120
tttctttctt agtcttctta tataagcgct atacgttcac attccatctc ctttggttat
                                                                       180
ctttccattt cttcaccgar cctctttgct ctctttttt atagctggtt cactcaaaat
                                                                       240
gtcttacttt gccatttttg aaatttattt tcattctttt atgtactgaa taaaatttaa
                                                                       300
aaatacttta tcatggtggg aggtacccgt gatgtccaaa taagtgttta tattaattgt
                                                                       360
tggggttttt ttgtttgtgt gttttttgaa aggttaagaa aatctcattc agaaagtaag
                                                                       420
ttgtttaaaa attctggacc aaatttacca cacatcaagc agatacttac caagttgttt
                                                                       480
ggtagacatt agcagtattt a
                                                                       501
<210> 56
<211> 541
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 432
<223> 99-15280-432 : polymorphic base C or T
<220>
<221> misc binding
<222> 409..455
<223> 99-15280-432.probe
```

```
<220>
<221> primer_bind
<222> 413..431
<223> 99-15280-432.mis
<220>
<221> primer bind
<222> 433..451
<223> 99-15280-432.mis complement
<220>
<221> primer bind
<222> 1..18
<223> 99-15280.pu
<220>
<221> primer bind
<222> 521..541
<223> 99-15280.rp complement
<400> 56
atgtccatcc atcttgccca gagagagttt ctacaacact tcctctgcaa gccctttccc
                                                                        60
tacttgcctc acctattgct ttcctctgtt acgttgtatt cccctcactg tttcttccaa
                                                                       120
catcttccca cctcagagca tggacacttg ctgctctttc tgtgtcatga tgctgctcac
                                                                       180
ttgtcccttt cttaatgtct cctccctgag ccaatcttct ccacccccac aacttacgca
                                                                       240
cacttacatg tcatattttc cttcatagcc tttaacacca tttgaaatga tatattttg
                                                                       300
attgctttta aaatttctct gtcccccac taaatataaa cttcaggatg gcaagaatgt
                                                                       360
agtccattat cttatttctc cagcctccat acttttaaga aaataaattt tggttgtata
                                                                       420
agccatccag tyagtggtac ttggttatag cacccctagc aaaagaatac aaaaaaggg
                                                                       480
agaatgtttg caatcatctg tttgaggcta ggaattccca gagagggaaa caaggagtaa
                                                                       540
                                                                       541
<210> 57
<211> 514
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 428
<223> 99-15353-428 : polymorphic base C or T
<220>
<221> misc binding
<222> 405..451
<223> 99-15353-428.probe
<220>
<221> primer_bind
<222> 409..427
<223> 99-15353-428.mis
<220>
<221> primer_bind
```



120

180

240

300

360

420

480

```
<222> 429..447
<223> 99-15353-428.mis complement
<220>
<221> primer_bind
<222> 1..18
<223> 99-15353.pu
<220>
<221> primer bind
<222> 495..514
<223> 99-15353.rp complement
<400> 57
tgggaatgga ggtagtagac gatgaggtct ccaccctctg actttgcaga gatgggcaag
gccaagtgtt ggaagggctt aaacacacac cggagtattc tgtgagaacc agtggatttc
agaggatggc aatgacacca cttgccttct gcctcaggag gataactgat ggccgtgtgt
gggatgcact ggagagcaag agctggcttg cagggagacc agctggatga ttttctttca
tttattttat tcattcaaca cacattcatc tggggttcac tctgtgccca acactgggca
tttccaaata gtccagatgg cagtaagcat ggttgtggca gtaggaatgg gaaggctggg
aggggtatga gaggcattac aaacgggaag tgggagtggc accccagaaa agtctagttt
aaggtgcyag tggatgtgt catgtgtgcg cgggggtgtc tagagggtgg cgggcagctg
gaaattgagg tcaagtgctt aaagaacaac tcgt
<210> 58
<211> 489
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 150
<223> 99-15355-150 : polymorphic base C or T
<220>
<221> misc_binding
<222> 127..173
<223> 99-15355-150.probe
<220>
<221> primer bind
<222> 131..149
<223> 99-15355-150.mis
<220>
<221> primer_bind
<222> 151..169
<223> 99-15355-150.mis complement
<220>
<221> primer bind
<222> 1..18
<223> 99-15355.pu
```

```
<220>
<221> primer_bind
<222> 471..489
<223> 99-15355.rp complement
<400> 58
taacttctcc gtctctcctt cttagcccat atgtcaataa tgactgaaag tattcatttc
                                                                        60
catctttaaa ctgcctattc cagccacctc ccacctccat ctctttcctt ctaagttttc
                                                                       120
ttcatcttct actttgggca aaaggaaaty gatgtgtcag acaggcctag ttttgaattc
                                                                       180
tggatctgct agcacttctc tgtgtgtcct tggttatatg atatagtctt aaaccttaat
                                                                       240
gttcttgcct gtaaaatggg gataataaaa acctcttaac agtggttgtt tcatgcagct
                                                                       300
ttcattacaa acttcctcat tcaaaatctt caatgatttc catttttcac aaaatgaaat
                                                                       360
tcaaaatttc tgtagattat tgagacaagt cccctactct tcacctaaat ttatctttta
                                                                       420
tttattctct catcattatc aacaactact aggctttgtt gccttgactc cagaggcaaa
                                                                       480
aatcttatc
                                                                       489
<210> 59
<211> 468
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 227
<223> 99-15685-227 : polymorphic base A or G
<220>
<221> misc_binding
<222> 204..250
<223> 99-15685-227.probe
<220>
<221> primer bind
<222> 208..226
<223> 99-15685-227.mis
<220>
<221> primer_bind
<222> 228..246
<223> 99-15685-227.mis complement
<220>
<221> primer_bind
<222> 1..18
<223> 99-15685.pu
<220>
<221> primer bind
<222> 449..468
<223> 99-15685.rp complement
<400> 59
aaacaaaggc acgcagagga taaggcatga gtccaaccag cagcatctcc ctcccgaatg
                                                                        60
agtacagaaa tgatcaatac tcgaagagaa aaagatgctt tcagtgtgct ttacctgaaa
                                                                       120
```





```
acttccttaa gcagcttcac tttattgtca ggatatcgct ttgtgtttgt atcatctaag
                                                                       180
aaagctcgcg catatgctag tgggccagca ttgacctaga caaagarcaa agattttcag
                                                                       240
ttccactagg aagaaaatca ccatgaccat ctgctcagtt tcagtttgca ggcactaaaa
                                                                       300
agcccgttcg cgtgagctac tcacaatccc tgccttccag gaacttaagc ccaaaaagaa
                                                                       360
accacaaagc tcactctgtt gcacaccact tgattccatg atctcagcca tcttcagggc
                                                                       420
acttgtgatg atggtttact ttatgtaaga agaaaccaat gcttggaa
                                                                       468
<210> 60
<211> 500
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 428
<223> 99-15695-428 : polymorphic base C or T
<220>
<221> misc binding
<222> 405..451
<223> 99-15695-428.probe
<220>
<221> primer bind
<222> 409..427
<223> 99-15695-428.mis
<220>
<221> primer bind
<222> 429..447
<223> 99-15695-428.mis complement
<220>
<221> primer bind
<222> 1..18
<223> 99-15695.pu
<220>
<221> primer bind
<222> 481..500
<223> 99-15695.rp complement
<400> 60
atcagccttt gtgaggagga ggccctgcct gctctcctcc tgagctgatg ggtcagtcac
                                                                       60
accaggacaa aggtctgccc ggggctgtgt gggttcctcc ttcctgagct gcacaccagc
                                                                      120
atctgctgaa caccttctgg agctcagctc agtgtctcgt ccagagacac tggttccctt
                                                                      180
ggcttctcag caactctcgg atctgggcct gggtctaacc tcagcggtgg tcttgcccat
                                                                      240
ttctagggcc tcacaattca gcctcatgtc ttcacctgtg gctcttttgc aaggctcaga
                                                                      300
aagetetagg gteagtteea gatgaeteee accageatge eagtaggage caccaeeeee
                                                                      360
tctcagccag cgccaccata ttccaggcaa attccaactg acacagactt caaggaacga
                                                                      420
ttgtagcygt tgttcttgct tcttccaaat ggaagagtgc attattgggg tcccttctag
                                                                      480
cacgcatttc attccccacc
                                                                      500
```

<210> 61

```
<211> 472
      <212> DNA
      <213> Homo sapiens
      <220>
      <221> allele
      <222> 310
      <223> 99-15703-310 : polymorphic base C or T
      <220>
     <221> misc binding
     <222> 287..333
     <223> 99-15703-310.probe
     <220>
     <221> primer bind
     <222> 291..309
     <223> 99-15703-310.mis
O
     <220>
⊭
     <221> primer_bind
     <222> 311..329
     <223> 99-15703-310.mis complement
     <220>
     <221> primer_bind
     <222> 1..18
     <223> 99-15703.pu
     <220>
     <221> primer_bind
     <222> 452..472
     <223> 99-15703.rp complement
     <400> 61
     agggctttgg gttataggcg ctgaatttct tctaaagcta acctgactct gatgctagaa
                                                                             60
     gagcccattt aaggaaagaa aaacactttt cattgctcga tcaaagttca tccattttgg
                                                                            120
     aaaagacatc aaaccaagtg tgtgacacca ggcacccata tccttcctct ttcccaccac
                                                                            180
     cccacccctg tcctcagggc agtgacagtg aagcctggtg caggtcccgc tgctgctttt
                                                                            240
     tgaagtggca catgctttat tttcttaaaa agaagtgaga gacaacctat gctacaggag
                                                                            300
     gctctgtgay gtttttctga agtacaaccc cttgctctgc cagggcagct gtaaagggtc
                                                                            360
     taaagagccc tgagaaagga gagaggattt gggaagccga ggaggcagag ggagaccaca
                                                                            420
     tagcacatgg agttctgaaa gggcccaagt ggagacagaa aacgagtcat gt
                                                                            472
     <210> 62
     <211> 470
     <212> DNA
     <213> Homo sapiens
     <220>
     <221> allele
     <222> 400
     <223> 99-15870-400 : polymorphic base A or G
```

<223> 99-16321-287.mis

```
<220>
<221> misc_binding
<222> 377..423
<223> 99-15870-400.probe
<220>
<221> primer_bind
<222> 381..399
<223> 99-15870-400.mis
<220>
<221> primer bind
<222> 401..419
<223> 99-15870-400.mis complement
<220>
<221> primer_bind
<222> 1..21
<223> 99-15870.pu
<220>
<221> primer bind
<222> 452..470
<223> 99-15870.rp complement
<400> 62
gctcaaatgt atcaaacaca gtttctgtgg tcaagttcct ctccttttct aaatttgctt
                                                                        60
agaggatete ataaaaegta aeteetetga caagggaaee attttageae caacaetgea
                                                                       120
aaagcttctg tgttcctaag ggaaagatcc tttcctgaat taaatttaac ctctttagta
                                                                       180
ctcccattta gccacctgat aaatccactt gagctatctt ttgggaagag agaggtatct
                                                                       240
gggaacaata acacttcctt tttgaacagt ttaataaagc tttgtgagat ttcaagatga
                                                                       300
aagataatgt gtaatgctga tagtgccctc caaggctctg cattcatgga tccaattacg
                                                                       360
ttttttgtca tggtaaaagc cacagtggat atattaaatr agagtgtggt ttaagaatga
                                                                       420
aggcccagga gtctggagat ctggtttcta aggctgactt cacttctgct
                                                                       470
<210> 63
<211> 469
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<223> 99-16321-287 : polymorphic base A or C
<220>
<221> misc binding
<222> 264..310
<223> 99-16321-287.probe
<220>
<221> primer_bind
<222> 268..286
```

```
<220>
<221> primer bind
<222> 288..306
<223> 99-16321-287.mis complement
<220>
<221> primer bind
<222> 1..20
<223> 99-16321.pu
<220>
<221> primer bind
<222> 451..469
<223> 99-16321.rp complement
<400> 63
ctttaggaat atcccttctg atttgaacaa cattttgcta tccaagttct gtctactttt
                                                                        60
ttaacaagtt cttgctccgt gtgtctcctt ttgcttgttc tcaagtaagg gagtaacagg
                                                                       120
gataaactcc cactccttgg taaatctttc tatcattttt ggaaatctca tccattgtag
                                                                       180
taaatgctct taaatcttca tcttcaggcc gtgacttcca tctagcctcc attcacgttt
                                                                       240
ccgggtttat gtctgcaatg agcattccgt ggctctacat agatgcmcca ccatacctag
                                                                       300
aacccatgta tcccaaactc aattctttct ttcccaggac attacttcct gcacttcctt
                                                                       360
agtctatcaa tggcactgtt attctcttga ccatctagac ttgaaatttt ggggtttgga
                                                                       420
ctcctcctgc tcccttgctt tatatgtaat cagacatcaa gtctcaatc
                                                                       469
<210> 64
<211> 544
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 194
<223> 99-16333-194 : polymorphic base A or G
<220>
<221> misc_binding
<222> 171..217
<223> 99-16333-194.probe
<220>
<221> primer bind
<222> 175..193
<223> 99-16333-194.mis
<220>
<221> primer bind
<222> 195..213
<223> 99-16333-194.mis complement
<220>
<221> primer_bind
<222> 1..19
```

```
<223> 99-16333.pu
<220>
<221> primer bind
<222> 524..544
<223> 99-16333.rp complement
<400> 64
atttaccccg tctgccttgc aatttcagga tcagtataca tcaaatcaag tgaacaaccc
agggaattct gccgttacct tttagaaaca gaataaatat taacagagct ttacttcttt
ccaccaagga ggactatatg ttaatacagt aatttacact ggaaaaaata taaatgaaag
ggtttagaac ctcrtaactt taaaaataac ataattcctc ctagaacatt cctttcactt
gtgattctca aagcactttg catttcccag ctattggcag ggctggaatt aggatcaaag
tatcactaaa tggtaggtga aataaatgtg aagctgattt tcaggagtac aggaatggag
tcatcaggcg actttaaagt taagaatctg ttggagcagc tgccaataaa tcaaggccca
aaggagaaag ttctttggaa accttgaaat attgtataca tttagataat tattgttgtt
gtcaatgtta acgaaaaaag caataaatca gggagatggc actgatgagt gaggagaaat
<210> 65
<211> 475
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 149
<223> 99-5873-159 : polymorphic base C or T
<220>
<221> misc_binding
<222> 126..172
<223> 99-5873-159.probe
<220>
<221> primer bind
<222> 130..148
<223> 99-5873-159.mis
<220>
<221> primer bind
<222> 150..168
<223> 99-5873-159.mis complement
<220>
<221> primer_bind
<222> 1..18
<223> 99-5873.pu
<220>
<221> primer bind
<222> 457..475
<223> 99-5873.rp complement
```

120

180

240

300

360

420

480

540



```
<220>
<221> misc_feature
<222> 409
<223> n=a, g, c or t
<400> 65
gcgtaacaat aagcagggtt agtcgccaca aaacttgaga taagaggaaa actaaaaaag
                                                                        60
tctaatgaaa tcagtagtct taaaaaagatg acatgatagg aagagaagtg ttaaaaaaga
                                                                       120
aaaaaaatag gtatgaaaga gagtaacaya taccggaaaa gggataaaat acatcctttg
                                                                       180
aaagaacaaa gagttattca aattgaattc ttaatgaatt acttaaacag cagattagat
                                                                       240
attgttaaaa agaggaatag ggaattaaat gatatatgtg atgatattac ctagtgtaac
                                                                       300
catcaaagat gtattgcaaa tgataaagaa aaaaatgctg ccatggcaat attaatatca
                                                                       360
taaaaatata ctttaagaag taaataaatg caactaggaa tagagaaans dvhatgaata
                                                                       420
ataatattta amaaavvgta taacaagtat acataagatg taatatccta aaccg
                                                                       475
<210> 66
<211> 511
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 49
<223> 99-5912-49 : polymorphic base A or G
<220>
<221> misc binding
<222> 26..72
<223> 99-5912-49.probe
<220>
<221> primer bind
<222> 30..48
<223> 99-5912-49.mis
<220>
<221> primer bind
<222> 50..68
<223> 99-5912-49.mis complement
<220>
<221> primer bind
<222> 11..31
<223> 99-5912.pu
<220>
<221> primer bind
<222> 494..511
<223> 99-5912.rp complement
<400> 66
aaatataata gtcaaatcat gttaccatta ggacacatta aaaatgtcra attaccttgg
                                                                       60
gaccttatat gaacatatta agataataat gatagtgttc agtgcaatat tcagatcaat
                                                                      120
agtttaaacc caaaatattt ataccttcag attagatgta tgcaaatgca ttgattcatg
                                                                      180
```



```
tgtcttttat ctgttgttta catttggaga aatatttgag aaatatttca aaatggaatt
                                                                        240
tatataaatt taaacacata atggttttat gtaaaaatat tgctaaatta cattttcccc
                                                                        300
ttaattetta tttettggaa aegtgeetta gtegetgaaa tatteataea ttaacaeaat
                                                                        360
gaaagaagtg aaccttacta ggctttgact atcaggtttg ctgttggttt ttgactattg
                                                                        420
tgaaactata gcctgatttc taaatcagga agaaacgtgt attgttgtta atatggacac
                                                                        480
atgacatatt tgtctgcctg acttttgatc c
                                                                        511
<210> 67
<211> 485
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 210
<223> 99-6012-220 : polymorphic base G or T
<220>
<221> misc_binding
<222> 187..233
<223> 99-6012-220.probe
<220>
<221> primer bind
<222> 191..209
<223> 99-6012-220.mis
<220>
<221> primer bind
<222> 211..229
<223> 99-6012-220.mis complement
<220>
<221> primer bind
<222> 1..19
<223> 99-6012.pu
<220>
<221> primer bind
\langle 222 \rangle \ 467...485
<223> 99-6012.rp complement
<400> 67
gtcttgactt gttttcctga gggtccaggt tgatttgcat gctcttgagg aaatatacac
                                                                        60
gtcttctcag ttttaataat tgactgacag ccctgtggtt tctcaggacc cagtgagctg
                                                                       120
ctgctcccag gtcagtctgc aaaggatgct ggttcccttg tggtctcatc aaggtgagga
                                                                       180
atttcctgat tttagagatt tctttatcck aattttgaag actttctttc acatttctag
                                                                       240
gcataaaaaa atgtacagca ctctactgct tgtttaacaa atggatagtg atatatctgc
                                                                       300
caacaaagac cacatggagt atttcattga ctatcagaga agtttcctcg aaaggcacca
                                                                       360
tacttagtgt tttatttcca tgagtgaagg aaaattagtt atttgaagta tttggctgtc
                                                                       420
tttagttgtt tctaaagtag tgctgatttt atatgcccat aatattcata tatacaccca
                                                                       480
ggata
                                                                       485
```

<210> 68

<211> 529 <212> DNA

<221> allele <222> 89

<222> 66..112

<222> 70..88

<222> 90..108

<222> 1..18 <223> 99-6080.pu

<221> primer_bind

<221> primer_bind <222> 509..529

<220>

<220>

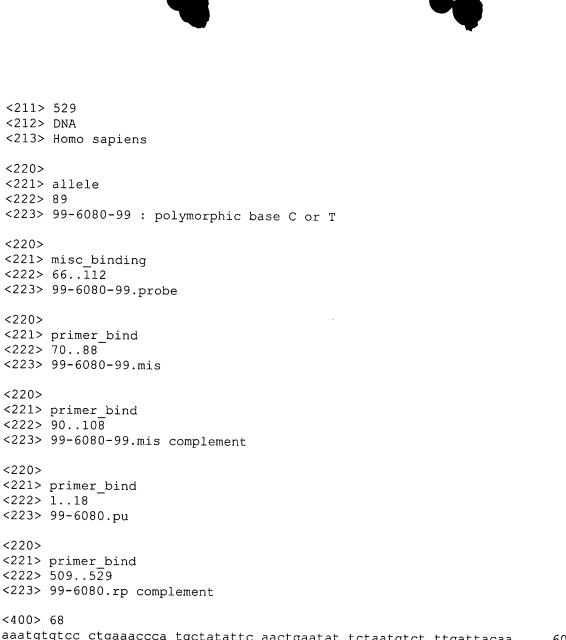
<220>

<220>

<220>

<220>

<400> 68



```
aaatgtgtcc ctgaaaccca tgctatattc aactgaatat tctaatgtct ttgattacaa
                                                                       60
agccatctct agcaatttaa tacaattayg aaatggaaaa gttggcaaat gcaaaacaat
                                                                      120
agctcgtgtt caaggtatgt ctttattagg ggaagtttat cgaaacagat gtttatgcta
                                                                      180
tttcctataa actagattct aaaatatttt attctataaa gatgtattga ctttatatga
                                                                      240
aaaaattatt gaaaaatcta caagatggtg aaactcttta gaactatatt tctattacaa
                                                                      300
gtttattttt aatttcaaaa atgtactgca taaatgcagc aaaaccttta ttgtcacata
                                                                      360
ttaaaacatg tacattattg tgtgcaaatt aaaatttcat taccttaaac caaaaagtga
                                                                      420
gttggccaga tagtaaataa tttaggctct aaggctgaaa agcgcttgta ttaattactc
                                                                      480
aactccacca ctattttgcc aaagcagtca cagacaatac gcattcaca
                                                                      529
<210> 69
<211> 489
```

```
<212> DNA
<213> Homo sapiens
<220>
<221> allele
<222> 156
<223> 99-7308-157 : polymorphic base C or T
```





```
<220>
<221> misc binding
<222> 133...179
<223> 99-7308-157.probe
<220>
<221> primer bind
<222> 137..155
<223> 99-7308-157.mis
<220>
<221> primer bind
<222> 157...175
<223> 99-7308-157.mis complement
<221> primer_bind
<222> 1..18
<223> 99-7308.pu
<220>
<221> primer_bind
<222> 469..489
<223> 99-7308.rp complement
<400> 69
tgtggtctgg atatggtgra ctgtccttca cacacagatg tgggaagcca tgatcatcag
                                                                        60
ttgcattatt cctgaggggc aatgcattcc agttacatag aaccagtttc tacgtttcag
                                                                       120
ggtatatgta ttcatggtga caaatttatt cacatyttaa gtaattttaa gtaattcaca
                                                                       180
ttttaagtaa ttttcctgaa tgtgcctcat tggcttctgt gcctcttcag aaaagatgaa
                                                                       240
ctaaacactg gcatatgtgt tcagatttca acattccgtt gttttcattg tggataattt
                                                                       300
ctgtcccata tttttgtgta aagttagaca ataaagtgtt aatattctgg cgtcggcaca
                                                                       360
ttttctttcc tgataaataa caattcacat atctttttaa aatatcagag aatatagtaa
                                                                       420
ccaatttcca attcttttt caccatgtat ctattggagt tttaaaatga ctaatactaa
                                                                       480
ggcaactat
                                                                       489
<210> 70
<211> 18
<212> DNA
<213> Artificial Sequence
<223> sequencing oligonucleotide PrimerPU
<400> 70
tgtaaaacga cggccagt
                                                                       18
<210> 71
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
```





<223> sequencing oligonucleotide PrimerRP

<400> 71 caggaaacag ctatgacc